#### STIC-Biotech/ChemLib

78693

From:

Zeman, Mary

Sent:

Friday, October 25, 2002 8:12 AM

To: Subject:

STIC-Biotech/ChemLib Sequence search 09/970966

Please run a <u>regular</u> search **and** an <u>oligo</u> search of the following sequences from 09/970966

SEQ ID NO: 199, 210, 211 and 214

please make the minimum window in the oligo searches 10 nucleotides.

please include interference files

Paper printout please.

Thank you, Mary K. Zeman Examiner, 1631 305-7133 CM1 12A17

mailbox: CM1 12D01

199- EFD 09/656,668

Point of Contact: Thomas G. Larson, Ph.D. 703-308-7309 CM1, Rm. 6 B 01

TYPE OF SEARGH:
NA Sequences: 4
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other

VENDOR/COST (where applic.)
STN:
DIALOG:
Questel/Orbit:
DRLink:
Lexis/Nexis:
Sequence Sys.:
WWW/Internet:
Other (specify):

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Maximum DB seq length: 2000000000
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Pred. No.

is the number of results predicted by chance to have a

REFERENCE AUTHORS TITLE

Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 369)
Xu,J. and Stolk,J.A.
Ovarian tumor sequences and methods of use therefor
Patent: WO 0118046-A 199 15-MAR-2001;

JOURNAL

SOURCE ORGANISM

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

DEFINITION ACCESSION VERSION KEYWORDS SOURCE	RESULT 1 AX093381 LOCUS		C 45	C 43		C 40	39	c 37	<u>ლ</u> ი		C 33	ω u u	c 30		27		0 C 25					c 18	16 17		) 14		c 11	_		c 7	ກປ	1.4	ω ι	<b>ب</b> د		т.	
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                                                                                    CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: National Institutes of Health Intramural
Sequencing Center (NISC),
Gaithersburg, Maryland;
                                                                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                     NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
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Submitted (05-NOV-2001) National Institutes of Health, Mammalian Submitted (MGC), Cancer Genomics Office, National Cancer Gene Collection (MGC), Cancer Genomics Office, National Ca
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This clone was selected for full length sequencing because it passed the following selection criteria: Hexamer frequency ORF analysis, Similarity but not identity to protein.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                  Homo sapiens
                                                            human.
                                                                                                      AX358762
                                                                                                                     Sequence 15 from Patent WO0193983.
                                                                                                                                                                AX358762
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zhang, L.-H. and Green, E.D.
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/protein_id="AAH17318.1"
/db_xref="GI:16878240"
/translation="MMVLGIAATFCGLFLLPGFALQIOCYQCEEFQLNNDCSSPEFIV
NCTVNVQDMCQKEVMEQSAGIMYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPLCNGPRPKKRGSSASALRPGLRTTILFLKLALFSAHC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="MGC:29643 IMAGE:3641660"
/tissue_type="Pancreas, adenocarcinoma"
/clone_lib="NIH_MGC_39"
/lab_host="DH10B-R"
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/db_xref="taxon:9606"
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Genentech
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                                                                                                                                                                                                                                                                                                          1524 bp
Sequence 15 from Patent W00208288.
AX362255
                                                                                                                                                                              Baker, K.P., Desnoyers, L., Gerritsen, M.E., Goddard, A., Godowski, P.J., Grimaldi, J.C., Gurney, A.L., Smith, V., Watanabe, C.K. and Wood, W.I.
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                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                               Unpublished
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                                                                            /organism="Homo sapiens"
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AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
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                                                                                                                                                                                                                                                                                              Submitted (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team); 2-6-7 Kazusa-Kamatari, Kisarazu, Chiba 292-0812, Japan (E-mail:genomics@hri.co.jp, Tel:81-438-52-3975, Fax:81-438-52-3986) NEDO human cDNA sequencing project supported by Ministry of Economy, Trade and Industry of Japan; cDNA full insert sequencing: Research Association for Biotechnology (RAB); cDNA library construction: Helix Research Institute (HRI) (supported by Japan Key Technology Center etc.); 5'- & 3'-end one pass sequencing: RAB, HRI, and Biotechnology Center, National Institute of Technology and Evaluation; clone selection for full insert sequencing: HRI and RAB, annotation: HRI and RAB.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kawakami,B., Sugiyama,A., Takemoto,M., Sugiyama,T., Irie,R., Otsuki,T., Sato,H., Wakamatsu,A., Ishii,S., Yamamoto,J., Isono Kawai-Hio,Y., Saito,K., Nishikawa,T., Kimura,K., Yamashita,H., Matsuo,K., Nakamura,Y., Sekine,M., Kikuchi,H., Kanda,K., Wagatsuma,M., Murakawa,K., Kanehori,K., Takahashi-Fijii,A.,
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oligo capping; fis (full insert sequence).
Homo sapiens alzheimer cortex cDNA to mRNA,
clone:BRALZ2001350.
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/clone="BRALIZ0001350"
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215..640
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                                                                                                                                                                                                                                                                               Secretory protein or membrane protein Patent: EP 1067182-A 203 10-JAN-2001;
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Sequence 203 from Patent EP1067182.
AX136281
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274. .771
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1 501 c 513 g 407 t
                                                                                                                                       /codon_start=
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                                                                                                                                                                                                             /organism="Homo sapiens"
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Query Match

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/db_xref="GI:17901492"
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/translation="MASPSLPGSDCSQIIDHSHVPEFEVATWIKITLILVYLIIFVMG
/translation="MASPSLPGSDCSQIIDHSHVELICHPEVATWIKITLILVYLIIFVMG
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SYTLSCKLHTFLFEACSYATLLHVLTLSFERYIAICHPFRYKAVSGPCQVKLLIGFVW
VTSALVALPLLFAMGTEYPLVNVPSHAGLTCNRSSTRHHEQPETSNMSICTNLSSRWT
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ARRQTIIFLRLIVVTLAVCWMPNQIRRIMAAAKPKHDWTRSYFRAYMILLPFSETFFY
LSSVINPLLYTVSSQQFRRVFVQVLCCRLSLQHANHEKRLRVHAHSTTDSARFVQRPL
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/db_xref="taxon:9606"
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          GCATCAGCTGCTGCAACACCCCTTTTGTAACGGGCCAAGG
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GCATCAGCTGCTACCACCCCTCTTTGTAACGGGCCAAGG
                                            GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT
                                                                                         AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 219
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Sequence 9
AX093191
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/db_xref="taxon:9606"
1, .396
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Sequence 478 from Patent
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The sequence of Homo sapiens
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MO 63108, USA
4 (bases 1 to 129676)
Waterston, R.H.
                                   Submitted (10-SEP-2000) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St.
                                                                   3 (bases 1 to 129676) Waterston, R.H.
                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 129676)
Sulston, J.E. and Waterston, R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Direct Submission

Submitted (09-JAN-2002) Department of Genetics, Washington
University, 4444 Forest Park Avenue, St. Louis, Missouri 63108,
On Aug 9, 2001 this sequence version replaced gi:14488388.

On Aug 9, 2001 this sequence Yearson Replaced Sequencing Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RP11-258B17 contains a single plasmid region from 1230 to 1239 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The clone sequenced to the left is RP11-1L22; the clone sequenced to the right is RP11-159N20, 2000 bp overlap. Actual start of thi clone is at base position 1 of RP11-258B17; actual end is at base position 60003 of RP11-159N20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mapping information for this clone was provided by Dr. John D. McPherson, Department of Genetics, Washington University, St. Louis MO. For additional information about the map position of this sequence, see http://genome.wustl.edu/gsc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NEIGHBORING SEQUENCE INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (http://bacpac.med.buffalo.edu)
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Contact: sapiens@watson.wustl.edu
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                                                                                            /rpt_family="AT_rich"
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JOURNAL REFERENCE
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                                                                                       2 (bases 1 to 147131)
VanBrunt, A. and Strowmatt, C.
The sequence of Homo sapiens BAC clone RP11-159N20
Unpublished (2001)
3 (bases 1 to 1777)
№ 4
            Submitted (28-SEP-1999) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St. MO 63108, USA
                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 147131)
Sulston, J. E. and Waterston, R.
Toward a complete human genome sequence
Genome Res. 8 (11), 1097-1108 (1998)
                                                           Direct Submission
                                                                      3 (bases 1 to Waterston, R.H.
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Homo sapiens BAC clone RP11-159N20
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                                                                                                                                     Submitted (01-MAR-2002) Department of Genetics, Washington University, 4444 Forest Park Avenue, St. Louis, Missouri 6. On Mar 1, 2002 this sequence version replaced gi:13435273.
                                                                                                                                                                                                                                                                                                                           Submitted (23-MAR-2001) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St. Louis,
                                                                                                                                                                                                                                      Direct
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                                                                                                                                                                                                                                                              Waterston, R
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                                                                                                                                                                                                                                                                                    (bases 1 to 147131)
                     Center: Washington University Genome Sequencing Center code: WUGSC
Genter code: Http://genome.wustl.edu/gsc
Contact: sapiens@watson.wustl.edu
Contact: Summary Statistics
Center project name: H_NH0159N20
                                                                                                                                                                                                                                      Submission
                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                        63108, USA
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NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.

all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality > 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by restriction digest. This sequence was finished as follows unless otherwise noted:

MAPPING INFORMATION:
Mapping information for this clone was provided by Dr. John D. McPherson, Department of Genetics, Washington University, St. MO. For additional information about the map position of this sequence, see http://genome.wustl.edu/gsc Louis

#### SOURCE INFORMATION:

The RPCI-11 human BAC library was made from the blood of one male donor, as described by Osoegawa, K., Woon, P.Y., Zhao, B., Frengen, E. Tateno, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong and coworkers at http://www.chori.org pBACe3.6

## NEIGHBORING SEQUENCE INFORMATION:

The clone sequenced to the right is RP11-458A7, 200 bp overlap; the clone sequenced to the right is RP11-258B17. Actual start of this clone is at base position 1 of RP11-159N20; actual end is at base position 147131 of RP11-159N20.

There clone are polymorphic base differences RP11-159N20 and RP11-258B17. 'n the overlap between the

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FEATURES
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                                                  /rpt_family="L1"
984. .1176
                                                                                        /rpt_family="MIR" 118. .1013
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991. .1176
                                                                                                                                                                                                /db_xref="taxon:9606"
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BG184021 (NID:g13705708)"
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/note="match to EST BG184021 (NID:g13705708)"

9743. .9895
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8373. .8683
                                                                                                                                                                                                                                                                                                                                                        /note="match to EST 15183. .15321
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9731. .9895
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16665. .16728
                                    16665
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1035. .1200
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/note="match to EST
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7811. .7881
                                                                                                                                                                                           note="match to EST BG187166 (NID:g13708853)"
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3348. .9395
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                                                   note-"match to EST BG704350 (NID:g13977603)"
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                                                                                      o/e=0.80, #CpGs=194)"
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Submitted (05-JUL-2002) Genome Sequencing Center, 4444 Forest Parkway, St. Louis, MO 63108, USA On Jul 5, 2002 this sequence version replaced gi:21426614.
                                                                        Submitted (14-JUN-2002) Genome Sequencing Center, 4444 Forest Park Parkway, St. Louis, MO 63108, USA 3 (bases 1 to 209885) McPherson,J.D. and Waterston,R.H.
                                                                                                                                               2 (bases 1 to 209885)
McPherson, J.D. and Waterston, R.H.
Direct Submission
                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 20985)
MCPherson, J.D. and Waterston, R.H.
The sequence of Mus musculus clone
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AC124493.2 GI:21699722
HTG; HTGS_PHASE1; HTGS_1
                                                                                                                                                                                                                                                                                                                                                                                                                                AC124493 209885 bp DNA line Mus musculus chromosome UNK clone RP23-462P13,
                                                         Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="similar to Bos taurus EST BE750540 (NID:g10164532)" 19498. .19795
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(NID:g11767217)"
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18931. .19137
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7. .19184
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           _family="AT_rich"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .19426
                                                                                                                                                                                                                                                                                                                                                         HTGS_DRAFT; HTGS_FULLTOP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 147131;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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13, WORKING
                                                                                                                                                                                                                                                                                                                                                                                                                                                 HTG 05-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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                                        Park
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Db 103249 GTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGC 103203
                                                                                                                                                                                                                                                                                         BASE COUNT
ORIGIN
                                                          RESULT 13
AB041649
    ACCESSION
                      DEFINITION
                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                             misc_feature
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                                                                                                                                                                      122 GTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGC 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequencing vector: M13; 0%
Sequencing vector: plasmid; 100%
Sequencing vector: plasmid; 100%
Chemistry: Dye-primer ET; 0% of reads
Chemistry: Dye-terminator Big Dye; 100% of reads
Assembly program: Phrap; version 0.990319
Consensus quality: 208484 bases at least 040
Consensus quality: 208911 bases at least 030
Consensus quality: 209285 bases at least 020
Insert size: 206000; agarose-fp
Insert size: 212112; sum-of-contigs
Quality coverage: 11.62 in 020 bases; sum-of-contigs
Quality coverage: 11.62 in 020 bases; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Center project name: M_BA0462P13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              web site:http://genome.wustl.edu/gsc/index.shtml
Contact: submissions@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Center: Washington University Genome Sequencing Center Center code: WUGSC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           * NOTE: This is a 'working draft' sequence. It currently consists of 5 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence.
Mus musculus brain cDNA, clone MNCb-0671. AB041649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      as soon as it is available and the accession number will be preserved.
                                                                                                                                                                                                                                                                                                           59789 a
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11432
28618
28718
49455
49555
107385
                                                                                                                                                                                                                Conservative
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/note="assembly_name:Contig11"
107485. .209885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1. .209885
                                                                                                                                                                                                                                                                                                                                                                                                                         /note="assembly_name:Contig9"
28718. .49454
                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="assembly_name:Contig8"
11432. .28617
                                                                                                                                                                                                                                                                                                         /note="assembly_name:Contig12"
44784 c 46164 g 58745 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:10090"
/chromosome="UNK"
                                                                                                                                                                                                                                                                                                                                                                                                     /note="assembly_name:Contig10"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="RP23-462P13"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11331: contig of 11331 bp in length
11431: gap of unknown length
28617: contig of 17186 bp in length
28717: gap of unknown length
49454: contig of 20737 bp in length
49554: gap of unknown length
107384: contig of 57830 bp in length
107484: gap of unknown length
209885: contig of 102401 bp in length.
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                                                                                                                                                                                                                                  12.7%;
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                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                    Score 47;
Pred. No.
                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                  DB 2; L
                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                    Length 209885;
                                                                                                                                                                                                                                                                                                               403 others
                                              linear
                                              ROD 30-JUN-2000
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                                                                                                                                                                                  В
                                                                                                                                                                                                                                                                                                                                                         BASE COUNT
              VERSION
                                  ACCESSION
                                                                    DEFINITION
                                                                                                      RESULT 14
AL512284/c
                                                                                                                                                                                                                                                                                                                                         ORIGIN
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                                                                                                                                                                                                                                                              38;
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              AL512284
AL512284.20 GI:18477310
                                                                                          AL512284
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FEATURES

Snoo

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VERSION
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                                                                                                                                                                                                                                                                       50 CAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGA 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AB041649.1 GI:7670499
fis (full insert sequence).
Mus musculus (strain:C57BL) adult female cDNA to mRNA,
clone_lib:Sugano mouse brain mncb clone:MNCb-0671.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ist strand cDNA was primed with an oligo(dT) primer lst strand cDNA was primed with an oligo(dT) primer lst strand cDNA was synthesized using specific 5' and 3' primers and amplified by PCR. The PCR product was digested with SfiI and size selection was performed to exclude fragments <1.5kb. The SfiI-digested PCR product was cloned into distinct DraIII sites of pmE18s-FL3. XhoI sites just outside the DraIII sites of pmE18s-FL3. XhoI sites just outside the DraIII sites of pmE18s-FL3. XhoI sites just outside the DraIII sites on be used to isolate the cDNA insert. Library was constructed by Sugano et al. (University of Tokyo, Institute of Medical Science). Custom primer used for sequencing (5' end primer [CTTCTGCTCTAAAAGCTGCG]; 3' end primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (13-APR-2000) Katsuyuki Hashimoto, National Institute of Infectious Diseases, Division of Genetic Resources; 23-1, Toyama 1-chome, Shinjuku-ku, Tokyo 162-8640, Japan (E-mail:khashi@nih.go.jp, URL:http://www.nih.go.jp/yoken/genebank/.Tel:81-3-5285-1111(ex.2120), Fax:81-3-5285-1181)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Osada,N., Kusuda,J., Tanuma,R., Ito,A., Hirata,M., Sugano,S. and Hashimoto,K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      made by oligo-capping method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   isolation of full-length cDNA clones from mouse brain cDNA library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A part of this sequence is re
Location/Qualifiers
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                       Human DNA sequence from clone RP11-248M22 on chromosome 10
complete sequence.
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lame: Sugano mouse brain mncb
                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /protein_id="BAA95101.1"
/protein_id="BAA95101.1"
/db_xref="GI:7670500"
/translation="MMVLGIAATFCGLFWLPGLALQIQCYQCEEFQLNNDCSSPEFIV
NCTVNVQDMCQKEVTEQSAGIMYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPLCNGPRPKKRGSSASAIRPGLLTTLLFFHLALCLAHC"
a 487 c 511 g 408 t
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/strain="C57BL"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone_lib="Sugano mouse brain mncb"
/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /codon_start=:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="unnamed protein product"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /sex="female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              clone="MNCb-0671"
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                                                                                                                                                                                                                                                                                                                                                                          10.3%;
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                                                                                                                                                                                                                                                                                                                                                                       Score 38; pred. No.
                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                              113920 bp
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                                                                                                                                                                                                                                                                                                                                                                             5e-11;
                                                                                                                                                                                                                                                                                                                                                                                                              DB 10;
                                                                     DNA
                                                                                                                                                                                                                                                                                                                                                                                                              Length 1797;
                                                                 linear
                                                                 PRI 31-JAN-2002
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В
                                                                                                                                    DEFINITION
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AUTHORS
TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      233 CAGTCCTTCTGCTCCCCAGGGA 254
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                               *** SEQUENCING IN PROGRESS**** 69 unordered pleces.
AC118794
                                                                                                                                                                                      AC118794
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This sequence was finished as follows unless otherwise noted: all regions were either double-stranded or sequenced with an alternate chemistry or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by at least one plasmid subclone or more than one Ml3 subclone; and the assembly was confirmed by restriction digest. The following abbreviations are used to associate primary accession numbers given in the feature table with their source databases: Em: EMBL; Sw:, surgepton. The convergence of the converg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         http://www.sanger.ac.uk/Projects/C_elegans/wormpep This sequence was generated from part of bacterial clone contigs of human chromosome 10, constructed by the Sanger Centre Chromosome 10 Mapping Group. Further information can be found at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               http://www.sanger.ac.uk/HGP/Chr10
RP11-248M22 is from the library RPCI-11.1 constructed by the group
of Pieter de Jong. For further details see
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (31-JAN-2002) Wellcome Trust Sanger Institute, Hinxton, Cambridgeshire, CB10 1SA, UK. E-mail enquiries: humquery@sanger.ac.uk Clone requests: clonerequest@sanger.ac.uk On Feb 1, 2002 this sequence version replaced gi:18476636. During sequence assembly data is compared from overlapping clones.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone="RP11-248M22"
/clone_lib="RPCI-11.1"
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                                                                                                                                                                                                                                                                                                                         Submitted (18-JUL-2002) Human Genome Sequencing Center, Depa. of Molecular and Human Genetics, Baylor College of Medicine, Baylor Plaza, Houston, TX 77330, USA
On Jul 14, 2002 this sequence version replaced gi:20387310.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (21-APR-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Direct Submission
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Direct Submission
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Mammalia; Eutheria; Rodentia; Sclurognathi; Muridae; Murinae;
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                                           Center project Information Center project name: GVHE Center clone name: CH230-179C24
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                                                                                                                                                      Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                  Center: Baylor College of Medicine Center code: BCM
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Summary Statistics
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TITLE

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NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 69 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.

This record will be updated with the finished sequence.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequencing vector: pUC18; L08752
Chemistry: Dye-terminator Big Dye; 90% of reads
Chemistry: Dye-primer Big Dye; 10% of reads
Chemistry: Dye-primer Big Dye; 10% of reads
Assembly program: Phrap; version 0.990399
Note:data AC066614 [Drafting center UWMSC] and AC066611
[Drafting center UWMSC] were added for finishing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Direct Submission
Submitted (10-UUL-2001) Multimegabase Sequencing Center, Institute for Systems Biology, 4225 Roosevelt Way NE, Suite 200, Seattle, WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (22-FEB-2000) Multimegabase Sequencing Center, University of Washington, PO BOX 357730, Seattle, WA 98195, USA 3 (bases 1 to 157042)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rowen,L., Madan,A., Qin,S., Abbasi,N., Baradarani,L., Birditt,B., Bloom,S., Dors,M., Dickhoff,R., Fleetwood,P., Harrison,G., James,R., Kaur,A., Madan,A., Owen,M.P., Ratcliffe,A., Shaffer,T.
                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98105, USA
On Jul 10, 2001 this sequence version replaced gi:13162487
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Direct Submission
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Sequencing of human chromosome 15 D15S146-D15S117 region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: leerowen@systemsbiology.org
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Web site: http://chroma.mbt.washington.edu/msg_www
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ----- Summary Statistics
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/note="overlap with RP11-348P16 AC066611
a 28493 c 28053 g 47690 t
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/note-"This clone overlaps RP11-8068 14.

RP11-348P16 AC066611. Data from overlapping BACs were
combined and the consensus sequence determined from
RP11-232J12 to the extent possible."
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143570. 157042
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FEATURES

Location/Qualifiers

Exon/Intron boundaries of identified genes were chosen if there were canonical splice junctions that maintained sequence continuity across the splice junctions.

Sequence similarities were identified using Powerblast by Jinghui The repeat regions shown were identified using RepeatMasker by Adrian Smit.

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COMMENT

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              On Dec 1, 2000 this sequence version replaced gi:11321701. Sequencing is completed to a minimum standard of double strand coverage with a minimum of 2 clones and 2 reads with no ambiguities or 2 chemistries with a minimum of 2 clones and 3 reads with no ambiguities. If the sequence quality does not meet this standard, it will be indicated in the annotation.
                                                                                                                              Submitted (01-MAR-2001) Human Genome Sequencing Center, Depart of Molecular and Human Genetics, Baylor College of Medicine, Baylor Plaza, Houston, TX 77030, USA On Dec 1, 2000 this sequence version replaced gi:11321701.
                                                                                                                                                                                                                                                                                             Submitted (05-DEC-2000) Human Genome Sequencing of Molecular and Human Genetics, Baylor College Baylor Plaza, Houston, TX 77030, USA 5 (bases 1 to 171146)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (01-DEC-2000) Human Genome Sequencing of Molecular and Human Genetics, Baylor College Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fernandez,C., Ferraguto,D., Forcum Tansey,J., Gill,R., Gorrell,J.H., Gunaratne,F., Haller,G., Hernandez,J., Hogues,M., Hosak,H., Hubur,J., Jackson,L., Jia,Y., Kelly,J., Kelly,S., Kovar,C., Liu,J., Liu,W., Loulseged,H., Lozado,R.J., Martin,R., Massey,E., McLeod,M.P., Mei,G., Moore,S., Morgan,M., Morrits,S., Massey,E., McLeod,A., Nguyen,R., Nguyen,N., Oguh,M., Parish,B., Perez,L., Reiter,D., Say,J., Shen,H., Vasquez,L., Watlington,S., Williamson,A., Wrensford,G., Zhou,X., Bouck,J., Hodgson,A., Muzny,D.M., Rives,M., Scherer,S., Sodergren,E., Weinstock,G., Muzny,D.M., Rives,M., Scherer,S., Sodergren,E., Weinstock,G., Muzny, K., and Gibbs,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     of Molecular and Human
Baylor Plaza, Houston,
4 (bases 1 to 171146)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (10-MAY-2000) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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Brooks,A., Brown,J.,
                                                                                                                                                                                                                                                                                                                                                  Center, Department of Medicine, One
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of Medicine, One
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TITLE

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                              DEFINITION
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Db 156695 CTCTGCCGGGTACCAGTCCT 156714
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180339 bp DNA linear PRI 12-JAN-
Homo sapiens chromosome 3 clone RP11-291P10, complete sequence.
AC099049 AC021722
AC099049.2 GI:18139443
                                                                                                                                                                                                l Similarity
20; Conserv
                                                                                                                                                                                            5.4%; Score 20; DB 10; Length 171146; llarity 100.0%; Pred. No. 3.9; Conservative 0; Mismatches 0; Indels 0;
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				JOURNAL			KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS
This sequence was finished as follows unless otherwise noted: all regions were either double-stranded or sequenced with an alternate chemistry or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by at least one plasmid subclone or more than one M13 subclone; and the assembly was confirmed by restriction digest.	th sequence assembly pro een reduced cted to have not generall available a	sembly program: Phrap; version or nsensus quality: 180276 bases at nsensus quality: 180338 bases at nsensus quality: 180339 bases at sert size: 180339; sum-of-contiguality coverage: 10.6x in Q20 bases alto per sequences: RPII-442C9 (UWGC:bc0425) AC02: RPII-59E22 (UWGC:bc0181) AC02	ton.ed  ion  ion  plo (b  80% c  40% c  40% c		98195, USA Zhou,Y., James,R.A., Rouse,G., n,P.K.A., Raymond,C. and Hauge	39) V., Raymond,C. and Hauger	HTG.  Homo sapiens.  Homo sapiens.  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  1 (bases 1 to 180339)  Kaul, R.K., Olson, M.V., Zhou, Y., James, R.A., Rouse, G., Wu, Z.,  Saepphimachak, C., Karen, P.K.A., Raymond, C. and Haugen, E.D.

in the table. There are no significant remaining discrepancies between the experimental and predicted values. Uniquely ordered fragments are separated by dashed lines.	
t remaining discrepancies d values. Uniquely ordered es.	

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SULT 19 025029 CUS FINITION CESSION RSION RSION GRANISM ORGANISM	308 <i>t</i> 178159 <i>t</i>	Duery Match Best Local Matches 2																									
ACO25029 ACO25029 Homo sapiens chromosome 3 cla 25 unordered pieces. ACO25029 ACO25029 ACO25029.14 GI:22203847 HTG; HTGS_PHASEL; HTGS_DRAFT human. Homo sapiens Eukaryota; Metazoa; Chordata Mammalia; Eutheria; Primates 1 (bases 1 to 190914)	AGGCCCAAGAAAAGGGGAAG               AGGCCCAAGAAAAGGGGAAG	11 Similarity 100.0%; 20; Conservative	i	<u></u>	1276		:	2859		2406	5229	27	27	4.4	8081	1	2591	4708	3139		4			169	845	ш.	847
chromos pieces. GI:2220 ASEI; HT etazoa; theria;		5.4%; 100.0% ative	6536	13421	1373	<800	<800	2932	963	2314	5278	2932	2932	4708	8247	3499	2606	4986	3247	800	5278	3719	4422	<800	843	1230	843
, ,, , , , , , , , , , , , , , , , , ,	327 178140	Score 20; Pred. No. 0; Mismatch			6226		-	615		1		8025	5512	2306			3572	10043	789			115	7	14		10	
DNA RP11-442C9,		9; ;e			6213	9956	2474	<800	<800	11368	1916	8039	5457	2319	2319	1449	3566	9956	<800	1566	<800	1078	<800	14076	3263	1078	3114
linear WORKING tebrata;		Length 180339;	958	-	4708		7	Д.	46	i on i		2041	30		26	31	2189	19	5084	7782			4165		1		1542
HTG 24-AUG-2002 DRAFT SEQUENCE, Euteleostomi;		Gaps 0;	981	<800	4606	<800	<800	1231	4606	5042	878	2052	3053	2052	2673	3257	2052	2052	5042	7809	1160	6296	4154	3257	5042	1347	1573
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		COMMENT	REFERENCE AUTHORS TITLE		TITLE JOURNAL	REFERENCE	TITLE																				AUTHORS

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Alsbrooks, S. L., Amaratunge, H. C., Are, J. R., Ayle, M., Banks, T., Alsbrooks, S. L., Amaratunge, H. C., Are, J. R., Ayle, M., Banks, T., Barbaria, J., Benton, J., Bimage, K., Blankenburg, K., Bonnin, D., Bouck, J., Bowle, S., Brieva, M., Brown, E., Brown, M., Bryant, N. P., Bouck, J., Bowle, S., Brieva, M., Brown, E., Brown, M., Bryant, N. P., Bucky, C., Burch, P., Burkett, C., Burrell, K. L., Byrd, N. C., Carron, T. F., Carter, M., Cavazos, S. R., Cheoko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C. D., Cox, C., Coyle, M. D., Dathorne, S. R., David, R., David, R., Dellaney, K. R., Delgado, O., Denn, A. L., Ding, Y., Dinh, H. H., Dellaney, K. R., Delgado, O., Denn, A. L., Ding, Y., Dinh, K. J., Denhart, C., Edgar, D., Edwards, C. C., Bhaj, C., Bscotto, M., Falls, T., Ferraguto, D., Flagg, N., Ford, J., Foster, P., Frantz, P., Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Gorrell, J. H., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K. J., Johnson, R., Havlak, P., Hawes, A., Hernandez, J., Homsi, F., Howard, S., Huber, J., Hulyk, S., Hume, J., Jackson, L. E., Jackson, B., Ja, Y., Johnson, R., Jolivet, S., Joudah, S., Karlsson, E., Kelly, S., Khan, U., King, L., Kwary, C., Hollins, B., Karlsson, E., Kelly, S., Khan, U., King, L., Kwary, C., Hollins, B., Karlsson, E., Kelly, S., Khan, U., King, L., Kwarth, J., Kovar, C., Kratovic, J., Kureshi, A., Landry, N., Lecier, A., Lucier, R., Luna, R., Ma, J., Louis, R., Martindale, A., Martinez, E., Maheshwari, M., Mapua, P., Martin, R., Martindale, A., Martinez, E., Maheshwari, M., Napua, P., Martin, R., Martindale, A., Martinez, E., Mung, G., Metzker, M., Moser, M., Pickens, R., Primus, E., Pull, L., Wolyen, N., Novers, S., Oguh, M., Okwuonu, G., Wetzker, M., Shoon, I., Rohley, H., Korney, J., Perez, L., Pickens, R., Primus, E., Pull, L., Wolles, M., Rohl, Y., Perez, L., Pickens, R., Primus, E., Pull, J., Novers, S., Savery, G., Scherer, S., Sortt, G., Shen, H., Shooshtari, N., Sisson, I., Thomas, N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (24-AUG-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Aug 11, 2002 this sequence version replaced g1:9438348.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (04-MAR-2000) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA 3 (bases 1 to 190914)
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Direct Submission
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Direct Submission
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Sequencing vector: M13;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 168264 bases at least Q40
Consensus quality: 176241 bases at least Q30
Consensus quality: 181136 bases at least Q20
Estimated insert size: 191206; sum-of-contigs estimation
Quality coverage: 3x in Q20 bases; sum-of-contigs estimation
                                                                                                                                                                                                                                                                                                                               Center project name: HAJK
Center clone name: RP11-442C9
Center clone name: Rtatistics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Web site: http://www.hgsc.bcm.tmc.edu/
Drafting Center Code: BCM
Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Center: Baylor College of Medicine
Center code: BCM
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NOTE: This is a "working draft" sequence. It currently consists of 25 contigs. The true order of the pieces is not known and their order in this sequence record is
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                                                                     Sequence
AX392969
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This sequence was finished as follows unless otherwise noted: all regions were either double-stranded or sequenced with an alternate chemistry or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by at least one plasmid subclone or more than one M13 subclone; and the assembly was confirmed by restriction digest. The following abbreviations are used to associate primary accession numbers given in the feature table with their source databases: Em:, EMBL; SW:, SMISSPROT; Tr:, TREMBL; Wp:, WORMPEP; Information on the WORMPEP database can be found at
                                                                                                                                                                                                                                                                                                                                   20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  On Apr 7, 2002 this sequence version replaced g1:15591682. During sequence assembly data is compared from overlapping clones. Where differences are found these are annotated as variations together with a note of the overlapping clone name. Note that the variation annotation may not be found in the sequence submission corresponding to the overlapping clone, as we submit sequences with only a small overlap as described above.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (04-APR-2002) Wellcome Trust Sanger Institute, Hinxton. Cambridgeshire, CB10 1SA, UK. E-mail enquiries: humquery@sanger.ac.uk Clone requests: clonerequest@sanger.ac.uk
                                            AX392969.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fttp://www.sanger.ac.uk/Projects/C_elegans/wormpep RP23-180B18 is
from the RPCI-23 Mouse PAC Library
constructed by the group of Pieter de Jong.
For further details see http://www.chori.org/bacpac/home.htm
human
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/db_xref="taxon:10090"
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AX392971/c
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 13 from Patent W00214358.
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HLRSLGALRLRHLAIASLEDQNFRRLPGLHLHEIDNWPLLEEVAAGSLRGLMLTSLSV 
THINITAVPAAALRHQAHLTCLNLSHNPISTVPRGSFRDLVRLRELHLAGALLAVVEP 
QAFIGRORIRLLNLSNNLLSTLEESTFHSVNTLETLRVDGNPLACDCRLLWIYQRRKT
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LCRAEGEPAPTVAWVTPQHRPVTATSAGRARVLPGGTLEIQDARPQDSGTYTCVASNA
GGNDTYFATLTVRPEPAANRTPGEAHNETLAALRAPLDLTTILVSTAMGCITFLGVVL
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/db_xref="taxon:9606"
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/db_xref="GI:19701017"
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/db_xref="taxon:9606"
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                   RLRVLRLRGNQLKLIPPGVFTRLDNLTLLDLSENKLVILLDYTFQDLHSLRRLEVGDN
DLVFVSRRAFAGLLALEELTLERCNLTALSGESLGHLRSLGALRLRHLAIASLEDQNF
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LTAVPDGIPAETRILELSRNRIRCLNPGDLAALPALEELDLSENAIAHYEPGAFANLP
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RRLPGLLHLEIDNWPLLEEVAAGSLRGLNLTSLSVTHTNITAVPAAALRHQAHLTCLN
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RESULT 24
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Sequence 9 from Patent WO0248361.
AX463719
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DLVFVSRRAFAGLLALEELTLERCNLTALSGESLGHLRSLGALRLRHLAIASLEDQNF
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EAHNETLAALRAPLDLTTILVSTAMGCITFLGVVLFCFVLLFVMSRGRGQHKNNFSVE
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/db_xref="taxon:9606"
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Eutheria;
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Kawasaki, K., Kubo, T. and Natori, S.
                                                                                                                                                                      D49515
D49515.1 GI:1902918
                                                        Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroldea; Dictyoptera; Blattaria; J
                                                                                                  Periplaneta americana fat body cDNA to mRNA, clone:PL25.Periplaneta americana
                                                                                                                                                                                                                                 American cockroach clone
                                                                                                                                                                                                                                                           D49515
                                         Blattidae; Periplaneta.
                                                                                                                                           lectin-related protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished
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Matsumoto_T.K., Pardo,J.M., Takeda,S., Bressan,R.A. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nicotiana tabacum
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SIWGAMFTESFYFKPYLNEKSKPKMYRDSNGVSGFDKSDLQLDVFMVQHDMENMYMWV
FKDRPENALGKMQLRS YMNGHSRQGERPFPESADRGFIRSHRVGRKHYRGLSNPQCVH
GIEVVPSPNLMVLDEEERKRWKELTGRDVNFTIPPEASDFSSWRNLPNTEFELEWPAP
PIKCKKLLNGSGLNLSTQPSNHSNGDAMDLLLPVNGKRKKDFSNGNBEEECYLPVNPP
TSYQIPDLEIHPNEPYWLNEFSGVMRDVYGPVTAAKSIYEDDEGYLIIISLPFVDLQR
VKVSWRNTLTHGIIKVSCLSTSRISFIKRQNRTFKLEASSSEHCPAGEFVREIPLSTR
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407 c 532 g 754 t
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/db_xref="GI:11611667"
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/db_xref="taxon:4097"
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                                                        Submitted (25-AUG-1998) Kenzo Sato, Tottori University Faculty of Medicine, Department of Molecular Biology; 86 Nishimachi, Yonago, Tottori 683-8503, Japan (E-mail:kensato@grape.med.tottori-u.ac.jp Tel:+81-859-34-8039, Fax:+81-859-34-8274)
                                                                                                                                                  Direct Submission
                                                                                                                                                                                                                                                                                                      Saito, K., Nakamura, T., Komoda, H., Hori, N., Adachi, K., Ito, K.
                                                                                                                                                                                  Saito, K., Nakamura, T., Komoda, H., Hori, N., Adachi, K., Ito, K.
                                                                                                                                                                                                                                Unpublished
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HNF-3G; hepatocyte nuc
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                                                                                                                                                                                                                                                                   Isolation and characterization of the rat hepatocyte nuclear
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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Presence of a lectin family in Periplaneta americana, revealed by analysis of cDNA clones of lectins related to Periplaneta lectin
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/organism="Rattus norvegicus"
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PLHVNVDHTAVLCERSDSVYIVATISAPPRSPSAGYELLPGLGYYKFHTDYKNWYDAR
KTCIQEGAHLAVINSETESKALLKLWLDHPKKENDWRNDWAHIGFHDHYTEGQFVTIF
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/product="lectan.related protein"
/protein_id="BAA18916.1"
/db_xref="GI:1902919"
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/db_xref="taxon:6978"
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19; Conserv
                                                                                                                                                                                                                                                                                                                                               Drive, Walnut Creek, CA 94598, USA

3 (bases 1 to 22752)

DOE Joint Genome Institute and Stanford Human Genome Center Direct Submission
                                                                                                                                   Quality: Phrap Quality >=40 100% of Sequence;
Estimated Total Number of Errors is 0.

MOTE: This insert is not the entire sequence of the clone. It is clipped at the overlaps with AC004152 and AC005258. The number of bases overlapped with AC004152 is 7297 and with AC005258 is 10523.

Location/Qualifiers
                                                                                                                                                                                                                                                                                    Submitted (01-MAR-2002) DOE Joint Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 94598, USA On Mar 1, 2002 this sequence version replaced gi:17646901. Draft Sequence Produced by DOE Joint Genome Institute
                                                                                                                                                                                                                                                                                                                                                                                                                   Direct Submission Submitted (13-DEC-2001) DOE Joint Genome Institute,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 22752)
DOE Joint Genome Institute and Stanford Human Genome Center.
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2996. .3914
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2695. .2699
2769. .2774
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/protein_id="BAA32534.1"
/db_xref="GI:4432776"
/ta_nslat.ton="MASGTPGLYASGLNSGRVPVERPKRSVPPGGPQPGRVGAQARGM
/tGSVKMEAHDLAEWSYYPEAGE"
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                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
/chromosome="19"
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                                             /clone="LLNL-FOS_25H9"
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Query Match

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Best Local Similarity
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1 (bases 1 to 37635)

Lamerdin,J.E., McCready,P.M., Skowronski,E., Adamson,A.W., Burkhart-Schultz,K., Gordon,L., Kyle,A., Ramirez,M., Stilwagen,S., Bhan,H., Velasco,N., Garnes,J., Danganan,L., Poundstone,P., Christensen,M., Georgescu,A., Anila,J., Liu,S., Attix,C., Duarte,S., Andreise,T., Trankheim,M., Amico-Keller,G., Coefield,J., Duarte,S., Andreise,T., Trankheim,M., Amico-Keller,G., Kronmiller,B., Arellano,A., Montgomery,M., Ow,D., Nolan,M., Trong,S., Kobayashi,A., Olsen,A.O.
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AC004152
AC004152.1 GI:2894631
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fosmid 37308 overlaps cosmid F20489 to the left; there is currently a 3.3 kb gap to the right between 37308 and cosmid R30783. Location Qualifiers
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Direct Submission
Submitted (18-FEB-1998) Joint Genome Institute, Lawrence Livermore Submitted (18-FEB-1998) Joint Genome Institute, CA 94551, USA National Laboratory, 7000 East Ave., Livermore, CA 94551, USA Map and sequence oriented from p telomere to centromere.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and Carrano,A.V.

Sequence analysis of a 3.5 Mb contig in human 19p13.3 containing a
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                                                                                                                                                                                                                                                                       /note="predicted exon, profiame: 1, quality: good, s complement(150. 227) /rpt_family="MIR" complement/11an
                                                                                 /rpt_family="AluSq/x"
2576. .2872
                                                                                                                                                        /rpt_family="AluSq/x"
complement(1845. 2140)
/rpt_family="Aluy"
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                                                                                                                                                                                                                                                   complement(1189. .1234)
/rpt_family="MIR"
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/clone="fos37308"
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/db_xref="taxon:9606"
complement(3551. .3848)
                                         /rpt_family="AluSx" 2993. .3293
                                                                                                                                                                                                                                  complement(1805.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /chromosome="19"
                                                                                                                                                                                                                                                                                                                                                                                                                        /rpt_family="AluJb"
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                        /rpt_family="Aluy"
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d, score: 59.000"
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/rpt_family="AluJ"
complement(10667..10921)
 complement(12479.
                                                                                                                                               DNA
                                                                                                                                                    /note="BLASTN similarity to D89870 (1.0.99, score: 2.1e-209; database searche
                                                                                                                                                                                                                                                                                     /rpt_family="Aluy"
complement/inn--
                                                                                                                                                                                                                                                                                                                                                                                                                             /rpt_family="Aluy"
9409. .9695
                                /rpt_family="MER20"
l2211. .12270
                                                                                                                                                                                                                                                       /rpt_family="AluJ"
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complement(5378. .5511)

/rpt_family="Aln.Th"
                                                                                                                                                                                                                                                                                                                               rpt_family="Alusg/x"
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complement(6285. .
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4230 ...4530
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7472. .7588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="predicted exon, program: grail2exons_human_1.3,
frame: 2, quality: excellent, score: 97.000"
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663. .6749
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3866. .4229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rpt_fami
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                                                                                                                                     or store: 2.1e-209; database searched: nt; Homo sapiens for antizyme, complete cds"
                                                                                                                                                                                          _family="AluJo"
9, .21272
                                                                             _family="AluSg"
              family-"Alusc"
                                                                                                           family="Alusx"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               family="Alusq"
ement(5207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          _family="L1MB7"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            . 8853
                                                                                                                                                                                                                                                                                                                                                          earcted exon, program: grail2exons_human_1.3,
quality: excellent, score: 76.000"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .y="AluSx"
.12772)
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                                                                                                                                                                    .10524); match:
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Matches

Conservative

0;

Mismatches

Gaps

35070 AAAAGGGGAAGTTCTGCCT 35088

317 AAAAGGGGAAGTTCTGCCT 335

Best Local Similarity

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Query Match
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AA317001 EST188650 HCC cell line (metastasis to liver in mouse) II Homo sapiens cDNA 5' end similar to similar to ornithine decarboxylase antizyme (1.200); 1008
identity.-AA308557 EST179376 HCC cell line (matastasis to liver in mouse) II Homo sapiens cDNA 5' end similar to similar to ornithine decarboxylase antizyme (1.200); 1008
identity.-(14748..14942) AA307682 EST178550 Colon carcinoma (HCC) cell line Homo sapiens CDNA 5' end similar to similar to ornithine decarboxylase antizyme (1.195); 1008 identity.-(14758..14942) L49059 HUM010907F Homo sapiens retinal fovea EST HFD010907 sequence.(221..37); 1008 identity.-Other EST matches: T53070, T47553, T27358, T19460"
                                                                  /note-DDS similarity to overlapping ESTs:
/note-DDS similarity to overlapping ESTs:
AA317001 EST188650 HCC cell line (metastasis to liver in mouse) II Homo sapiens cDNA 5' end similar to similar to ornithine decarboxylase antizyme (201. 346); 99% identity.-AA308557 EST179376 HCC cell line (matastasis to liver in mouse) II Homo sapiens cDNA 5' end similar to similar to ornithine decarboxylase antizyme (201. 346); 100% identity.-AA307682 EST178550 Colon carcinoma (HCC) cell line Homo sapiens cDNA 5' end similar to similar to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="translation of first 68 residues of ODC antizyme; /note="translation of first 68 residues of ODC antizyme; Ornithine decarboxylase antizyme is likely expressed through translational frameshifting which results in the products of ORF1 and ORF2 being synthesized as a single polypeptide. In rat (D10706) the mRNA sequence at the junction of the two ORF5, UCC UGA U (253. .259, shown as ORF1 codons), is decoded as Ser-Asp through +1
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complement(14613...14683)
/rpt_family="CC_rich"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SSSESSRVSLHCCSNPGPGPRWCS"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /product="ODC antizyme ORF1"
/protein_id="AACO2802.1"
/db_xref="G1:2894632"
/tzanslation="MYKSSLORILNSHCFAREKEGDKPSATIHASRTMPLLSLHSRGG
                                                                                                                                                                                                                                                                                                                                                                                  /gene="OAZ"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="ornithine decarboxylase antizyme; 98% identical to
(U09202) ornithine decarboxylase antizyme [Homo sapiens]."
join(14797. .14942,16583. .>16640)
/gene="OAZ"
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/rpt_family="AT_rich"
complement(14430..14
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complement(12872. ...
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5.1%; Score 19; DB 9; Length 37635; 100.0%; Pred. No. 16;
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                                                                                                                                                                                                                                                                                     COMMENT
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TITLE
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JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                    JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HTG; HTGS_PHASE1
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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                                                                                                                                                                                                                                                                                     of Molecular and Human Genetics, Baylor College
Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                         Submitted (19-JUL-2002) Human Genome Sequencing Center, Department
of Molecular and Human Genetics, Baylor College of Medicine, One
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wu,C., Wu,Y., Wu,Y.F., Zhou,J., Weinstock,G. and Gibbs,R.
                                                                                                                                                                                                                                                                                                                                                                                                                   Direct Submission
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                                                                                                            web site: http://www.hgsc.bcm.tmc.edu/
Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (bases 1 to 142329)
                                                         Center project Information Center project name: GZNK Center clone name: CH230-432K14
                                                                                                                                                                                                                                                Center: Baylor College of Medicine
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vector:
                       Summary
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                                 Statistics
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*** SEQUENCING IN PROGRESS
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* NOTE: Estimated insert size may differ from sequence length

* (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

* NOTE: This is a 'working draft' sequence. It currently

* consists of 45 contigs. The true order of the pieces

* is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap: version 0.990329 Consensus quality: 103440 bases at least Q40 Consensus quality: 1108530 bases at least Q30 Consensus quality: 112007 bases at least Q20
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Homo sapiens BAC clone RP11-524B5
AC105316
AC105316.5 GI:20522215
                      Unpublished
                                        The
                                                                                       Sulston, J.E. and Waterston, R.
Toward a complete human genome sequence genome Res. 8 (11), 1097-1108 (1998)
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                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
        Waterston, R.H.
                                                Buatsi, D., Haakenson, W.,
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/db_xref="taxon:10116"
/clone="CH230-432K14"
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99586: contig of 5542 bg

99686: gap of unknown le

106051: contig of 6365 bg

106151: gap of unknown le

112416: contig of 6265 bg

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123175: contig of 10659 l

133279: gap of unknown le

132799: contig of 9524 bg

132899: gap of unknown le

132899: contig of 95430 bg
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5 (bases
                                                                                                             Submitted (29-MAY-2002) Department of Genetics, Washington University, 4444 Forest Park Avenue, St. Louis, Missouri 63108, USA On May 10, 2002 this sequence version replaced gi:20340527.
                                                                                                                                                                                        Waterston,
Direct Sub
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Direct Submission
Submitted (10-MAY-2002) Genome
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Submitted (30-APR-2002) Genome
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Center project name: H_NH0524B05
                                     Contact: sapiens@watson.wustl.edu
                                                      Center code: WUGSC
Web site: http://genome.wustl.edu/gsc
                                                                                        Center: Washington University Genome Sequencing Center
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NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions

This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by restriction digest.

#### MAPPING INFORMATION:

Mapping information for this clone was provided by Dr. John D. McPherson, Department of Genetics, Washington University, St. MO. For additional information about the map position of this sequence, see http://genome.wustl.edu/gsc ohn D. St. Louis

### SOURCE INFORMATION:

The RPCI-11 human BAC library was made from the blood of one male donor, as described by Oscegawa, K., Woon, P. J., Zhao, B., Frengen, E. Tateno, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from processing the construction of the construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from processing the construction of the construction Research Genetics, Inc. (http://www.reand coworkers at http://www.chori.org coworkers at OR: pBACe3.6 (http://www.resgen.com) or Pieter de

# The clone sequenced to the left is RP11-620C21, 2000 bp overlap; the clone sequenced to the right is RP11-138A33, 2000 bp overlap Actual start of this clone is at base position 70882 of RP11-620C21; actual end is at base position 47389 of RP11-138A23 Location/Qualifiers

NEIGHBORING SEQUENCE INFORMATION:

FEATURES

repeat\_region source /organism="Homo sapiens"
/db\_xxef="taxon:9606"
/chromosome="4" /clone\_lib="RPCI-11" 138. .322 /clone="RP11-524B5" .14857

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Submitted (11-NOV-2000) Genome Sequencing Center, Washington
University School of Medicine, 4444 Forest Park Parkway, St. Louis,
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Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalla; Eutherla; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 154693)

Sulston, J. E. and Waterston, R.

Toward a complete human genome sequence
Genome Res. 8 (11), 1097-1108 (1998)
                                                                                                                                                                                                                                                                                                                                               154693 bp DNA linear PRI 09-JA
Homo sapiens BAC clone RP11-215J17 from 4, complete sequence.
AC084750
AC084750.3 GI:17155079
Submitted (30-NOV-2001) Genome Sequencing Center, Washington
                  MO 63108, USA
4 (bases 1 to 154693)
Waterston, R.H.
Direct Submission
                                                                                                                                        2 (bases 1 to 154693)
Isak,A. and Kozlowicz,A.
The sequence of Homo sapiens BAC clone RP11-215J17
Unpublished (2001)
                                                                                                              3 (bases 1 to 154693) Waterston, R.H.
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42742. 42779
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42561. .42741
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40222. .40250
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34556. .34879
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Submitted (03-JAN-2002) Genome Sequencing Center, Washington
University School of Medicine, 4444 Forest Park Parkway, St. Louis,
MO 63108, USA
6 (bases 1 to 154693)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              University School of Medicine, 4444 Forest Park Parkway, St. Louis, MO 63108, USA 5 (bases 1 to 154693) Waterston, R. H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Direct Submission
Submitted (09-JAN-2002) Department of Genetics, Washington Submitted, 4444 Forest Park Avenue, St. Louis, Missouri 63108, On Nov 30, 2001 this sequence version replaced gi:16799050.
                                                                                                                                                                                                                                                            Data from AC091490 was used to finish this clone, AC084750 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                    The clone sequenced to the left is RP11-362M19, 2000 bp overlap; the clone sequenced to the right is RP11-509110. Actual start of this clone is at base position 151949 of RP11-362M19; actual end is at base position 154693 of RP11-215J17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          The RPCI-11 human BAC library was made from the blood of one male donor, as described by Osoegawa, K., Woon, P.Y., Zhao, B., Frenger, E., Tateno, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong and coworkers at the Roswell Park Cancer Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MAPPING INFORMATION:
Mapping information for this clone was provided by Dr. John D.
Mapping information for this clone was provided by Dr. John D.
McPherson, Department of Genetics, Washington University, St. Louis
MO. For additional information about the map position of this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Waterston, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOURCE INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            restriction digest.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This sequence was finished as follows unless otherwise noted:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       between neighboring data submissions.
                                                                                                                                                                                                                                                                                                                                                                                                     NEIGHBORING SEQUENCE INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                     (http://bacpac.med.buffalo.edu)
VECTOR: pBACe3.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Center project name: H_NH0215J17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: sapiens@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Center: Washington University Genome Sequencing Center Center code: WUGSC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Web site: http://genome.wustl.edu/gsc
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                /rpt_family="AT_rich" 6208. .6327
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4574 . .4607
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/rpt_family="MaLR"
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/chromosome="4"
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                                                                                                                                                             /map="4"
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13902. .14148
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               Alsbrooks, S.L., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbrooks, S.L., Amaratunge, K., Blankenburg, K., Bonnin, D., Bouck, J., Benton, J., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Bouck, J., Bowle, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Buhay, C., Burch, P., Burkett, C., Burch, K.L., Byrd, N.C., Carron, T.F., Carter, M., Cavazos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, G., Chen, R., Chowdhry, I., Christopoulos, C., Coyle, M.D., Dathorne, S.R., David, R., Davila, M.L., Davis, C., Coyle, M.D., Dathorne, S.R., David, R., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Delaney, K.R., Delgado, D., Edwards, C., Elhaj, C., Escotto, M., Falls, T., Ferraguto, D., Edwards, C., C., Elhaj, C., Escotto, M., Falls, T., Ferraguto, D., Flagg, N., Ford, J., Foster, P., Frantz, P., Garls, A., Garner, T., Garza, N., Gill, R., Gobisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Gobisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Harris, C., Harris, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J., Homsi, F., Howard, S., Huber, J., Huber, J., Jackson, L.E., Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S., Kratovic, J., Kureshi, A., Landry, N., Leal, B., Lewis, L.C., Lewis, L., Lichtarge, O., Lieu, C., Liu, J., Liu, W., Loulseged, H., Lozado, R.J., Lucler, A., Lucler, R., Luna, R., Ma, J., Mahachwari M. Mannia, D. Martin, B. Marti
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Maheshwari,M., Mapua,P., Martin,R., Martindale,A.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (13-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 12, 2002 this sequence version replaced g1:18701823.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (14-JAN-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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                                                                                                                                                                                                                                                                                                                                                     * arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Direct Submission
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Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 88710 bases at least Q40
Consensus quality: 93751 bases at least Q30
Consensus quality: 97432 bases at least Q20
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Center clone name: CH230-119A22
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Contact: hgsc-help@bcm.tmc.edu
------ Project Information
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* is believed the gaps between comments of the sequence as soon as it is available and the accession number will be preserved.

* the accession number will be preserved.

* 1 2249: contig of 2249 bp in length

* 2250 2349: gap of unknown length
                                                                                                                                                                                                                                                              OK 73019, USA
On Aug 1, 2002 this sequence version replaced g1:22002191.
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19; Conser
                                                                                                                                                    * NOTE: This is a 'working draft' sequence. It currently * consists of 6 contigs. Gaps between the contigs * are represented as runs of N. The order of the pieces
                                                                                                                                                                                                                  The University Of Oklahoma Center code: UOKNOR
                                                                                                                                                                                                                                                                                                   Direct Submission
Submitted (16-AUG-2002) Department Of Chemistry And Biochemistry, The University Of Oklahoma, 620 Parrington Oval, Room 208, Norman
                                                                                                                                                                                                                                                                                                                                                                                        Submitted (28-FEB-2002) Department Of Chemistry And Biochemistry, The University Of Oklahoma, 620 Parrington Oval, Room 208, Norman,
                                                                                                                                                                                                                                            Center: Department Of Chemistry And Biochemistry
                                                                                                                                                                                                                                                                                                                                                                               OK 73019, USA
                                                                                                                                                                                                                                                                                                                                                                                                                       Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished
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Mus musculus BAC Clone rp23-46k8
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156128 bp DNA linear HTG 16-AUG-2002
Mus musculus chromosome 17 clone rp23-46k8 strain C57BL/6J, WORKING
DRAFT SEQUENCE, 6 ordered pieces.
                                                                                                                                                                                                                                                                                                                                                 Milam,J.
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                                                                                                                           is believed to be correct as given, however the sizes of the gaps between them are based on estimates that have
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                                                                                                                                                                                                           Submitted (26-JUN-2001) Sanger Centre, Hinxton, Cambridgeshire, CB10 18A, UK. E-mail enquiries: humquery@sanger.ac.uk Clone requests: clonerequest@sanger.ac.uk
On Jun 27, 2001 this sequence version replaced gi:14272267.
During sequence assembly data is compared from overlapping clones.
Where differences are found these are annotated as variations together with a note of the overlapping clone name. Note that the variation annotation may not be found in the sequence submission corresponding to the overlapping clone, as we submit sequences with only a small overlap as described above.
This sequence was finished as follows unless otherwise noted: all regions were either double-stranded or sequenced with an alternate chemistry or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by a tleast one plasmid subclone or more than one M13 subclone; and the assembly was confirmed by restriction digest. The following abbreviations are used to associate primary accession numbers given in the feature table with their source databases: Em: EMBL; Sw:, SWISSPROT; TI:, TREMBL; Wp:, WORMPED; Information on the WORMPED than all property of the sequence of the content of the wormpen This sequence.
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AL354709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AL354709.15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human DNA sequence from clone
                          http://www.sanger.ac.uk/HGP/Chr9 RP11-145E5 is from the library RPCI-11.1 constructed by the group of Pieter de Jong. For further details see
                                                                                                                                  chromosome 9, constructed by the Sanger Centre
                                                                                                                                                                http://www.sanger.ac.uk/Projects/C_elegans/wormpep This sequence was generated from part of bacterial clone contigs of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Laird
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
http://www.chori.org/bacpac/home.htm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 45867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (bases 1 to 157533)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submission
                                                                                                           Further information can be found at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1. .156128
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/clone_lib="RPCI - 23 Female (C57BL/6J) Mouse BAC Library"
33372 c 32918 g 43471 t 500 others
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5.1%; Score 19;
100.0%; Pred. No.
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                                                                                                                                        Chromosome 9 Mapping
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Best Local
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Bouck, J., Boule, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P.,
Buhay, C., Burch, P., Burkett, C., Burrell, K.L., Byrd, N.C.,
Carron, T.F., Carter, M., Cavazos, S.R., Chacko, J., Chavez, D.,
Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C.,
Chen, G., Chen, R., Chen, D., Dathorne, S.R., David, R.,
Davila, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A.,
Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H. H.,
Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J.,
Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M.,
Falls, T., Ferraguto, D., Flagg, N., Ford, J., Foster, P., Frantz, P.,
Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R.,
Gorrell, J.H., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K.,
Harris, C., Harris, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J.,
Hernandez, O., Hodgson, A., Hogues, M., Holloway, C., Hollins, B.,
Hemsi, F., Howard, S., Huber, J., Hulyk, S., Hume, J., Jackson, L.E.,
Jacobson, E., Kelly, S., Khan, U., King, L., Korvah, J., Kovar, C.,
Karlsson, E., Kelly, S., Khan, U., King, L., Korvah, J., Kovar, C.,
Karlstovic, J., Kureshi, A., Landry, N., Leal, B., Lewis, L., C., Lewis, L.,
Li, J., Li, X., Lucler, A., Lucler, R., Luna, R., Ma, J.,
Massey, E., Mawhiney, E., McLeod, M.P., Meadorr, M., Mel, G., Metzker, M.,
Miner, G., Miner, Z., Mitchell, T., Mohabbat, K., Morgan, M., Morris, S.,
Moser, M., Neel, D., Newtson, N., Nguyen, A., Nguyen, N.,
Oviedo, R., Pace, A., Pace, A., Pavton, B., Peerv. J., Parez., I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IMPORTANT: This sequence is not the entire insert of clone RP11-145E5 It may be shorter because we sequence overlapping sections only once, except for a 100 base overlapping. The true left end of clone RP11-145E5 is at 1 in this sequence. The true right end of clone RP11-408N14 is at 157434 in this sequence. The true right end of clone RP11-149I2 is at 48466 in this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rattus norvegicus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rattus norvegicus clone CH230-354C23, ***, 59 unordered pieces.
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restriction digest data."
a 29239 c 31401 g 51244 t
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/db_xref="taxon:9606"
/chromosome="9"
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/clone_lib="RPCI-11.1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (18-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 14, 2002 this sequence version replaced gi:20467689.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (26-MAR-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA 3 (bases 1 to 163041)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Worley, K.C
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Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a "working draft" sequence. It currently consists of 59 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.

This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                            as soon as it is available and the accession number will be preserved.  
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap: version 0.990329
Consensus quality: 116366 bases at least Q30
Consensus quality: 120870 bases at least Q30
Consensus quality: 124554 bases at least Q30
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Center code: BCM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (25-APR-2000) Human Genomic Center, Institute of Genetics, Chinese Academy of Sciences, Datun Road, Beijing, Beijing 100101, P.R.China
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished 2 (bases 1 to 165873)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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HTG; HTGS.
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117935: contig of 4433 b
118035: gap of unknown 1
122534: contig of 4499 b
122634: gap of unknown 1
127202: contig of 4568 b
127302: gap of unknown 1
133412: contig of 5110 b
133512: gap of unknown 1
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of 3679 bp in length
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19; Conserv
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Db 65260 TCAGAAAGAAGTGATGGAG 65278
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Center project name: 1% project
Center clone name: RP11-34405
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                                                                                                                                                                                                                  HOMO SAPIENS Chromosome UNK clone RP11-289E6, WORKING DRAFT SEQUENCE, 31 unordered pieces.
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                    Eukaryota; Metazoa: Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 177070)
Waterston, R.H.
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AC018875.7 GI:12963043
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                                                                                                                                           Homo sapiens.
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The sequence of Homo sapiens clone
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Waterston,R.H.
Direct Submission
Submitted (21-DEC-1999) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St. MO 63108, USA
On Feb 18, 2001 this some
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chemistry: Dye-primer ET; 86% of reads chemistry: Dye-terminator Big Dye; 14% of reads Assembly program: Phrap; version 0.990319 Consensus quality: 163528 bases at least 040 Consensus quality: 169714 bases at least 030 Consensus quality: 173054 bases at least 020 Consensus quality: 173054 bases at least 020
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Insert size: 176607; sum-of-contigs
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Sequencing vector: plasmid; 14%
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39649. .41067
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25939. .27145
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109291: contig of 30138 bp in length
109391: gap of unknown length
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Sequencing vector: MIJ9: 100%
Sequencing vector: plasmid: 0%
Sequencing vector: plasmid: 0%
Chemistry: Dye-primer ET: 100% of reads
Chemistry: Dye-terminator Big Dye: 0% of reads
Assembly program: Phrap; version 0.990319
Consensus quality: 154559 bases at least Q40
Consensus quality: 166430 bases at least Q30
Consensus quality: 166430 bases at least Q20
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Submitted (15-CAN-2000) Genome Sequencing Center, Washington
University School of Medicine, 4444 Forest Park Parkway, St.
MO 63108, USA
On May 12, 2000 this sequence version replaced gi:7233676.
                                                                                                                                                                                                                          Center: Washington University Genome Sequencing Center Center code: WUGSC
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1 (bases 1 to 177426)
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                                                                                                                                                                   Center project name: H_NH0567M21
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61888. .65440
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Insert size: 189000; agarose-fp
Insert size: 174526; sum-of-contigs
Quality coverage: 2.77 in Q20 bases; agarose-fp
Quality coverage: 3.06 in Q20 bases; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                * NOTE: This is a 'working draft' sequence. It currently consists of 30 contids. The true order of the pieces is not known and their order in this sequence record is are rapitrary. Gaps between the contids are represented as runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           as soon as it is available and the accession number will be preserved.
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                                                          66090: gap of u
72519: contig
72519: gap of u
79568: contig
79468: gap of u
88731: gap of u
88731: gap of u
98171: contig
98271: gap of u
108874: gap of u
128856: contig
          133932: gap of unknown length 151337: contig of 17405 bp in 151437: gap of unknown length
                                        120956: gap of 133832: contig
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                                        f unknown length
g of 11882 bp in
f unknown length
g of 12876 bp in
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of 2845
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of 1877
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of 1470
                                                                                  g of 9440 bp in 1
f unknown length
g of 10603 bp in
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of 2649
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of 2823
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of 6749
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     Conservative
                                                                                      /note="assembly_name:Contig40"
151438. 177426
/note="assembly_name:Contig41"
38792 c 37341 g 49554 t
                                                                                                                                                                                                                                                                          /note="assembly_name:Contig36"
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66091 72519
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133933. .151337
                                                                                                                                                                                                                                                                                                                     /note="assembly_name:Contig35"
                                                                                                                                                                                                                                                                                                                                                             vector_side:left"
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120957. .133832
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/chromosome="11"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note="assembly_name:Contig16"
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                  5.1%; Score 19; DB 2; 100.0%; Pred. No. 16;
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AC112477
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Douthwaite, K. J., Delgado, O., Denn, A. L., Ding, Y., Dinh, H.,
Douthwaite, K. J., Draper, H., Dugan-Rocha, S., Durbin, K. J.,
Earnhart, C., Edgar, D., Edwards, C. C., Elhaj, C., Escotto, M.,
Falls, T., Ferraguto, D., Flagg, M., Ford, J., Foster, P., Frantz, P.,
Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R.,
Gorrell, J. H., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K.,
Harris, C., Harris, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J.,
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Massey, E., Martin, R., Martin, R., Martin, R., Ma, J.,
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Nguyen, N., Nickerson, E., Nochenkwo, S., Oguh, M., Okwuonu, G.,
Oragunye, N., Oviedo, R., Pace, A., Payton, B., Peery, J., Perez, L.,
Peters, L., Pickens, R., Primus, E., Pu, L.L., Quiles, M., Ren, Y.,
Rives, M., Rojas, A., Rojubokan, I., Shooshtari, N., Sisson, I.,
Sodergran, E., Sonaike, T., Sparks, A., Stanley, H., Stone, H.,
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Wall, M., Shooshtari, N., Wang, O.,
Wang, S., Watch, K., Palifedd, B., Wang, O.,
Wang, S., Watch, R., Wac
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Submitted (17-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One
                                                                                                                                                                                                                                                                                                       Submitted (21-FEB-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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Direct Submission
                                                                                                                                             Direct Submission
                                                                                                                                                                                                  Worley, K.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rattus norvegicus
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AC112477.3 GI:21745567
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***, 63 unordered pieces.
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COMMENT

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Baylor Plaza, Houston, TX 77030, USA on Jul 14, 2002 this sequence version replaced gi:20303368.
                                                                                                                                                                                                                                                                                                                                                                                                                                              NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank.draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 63 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as seas soon as it is available and the accession number will
                                                                                                                                                                                                                                                                                                                                                                                                                                     as soon as it is available be preserved.
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Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 138102 bases at least Q40
Consensus quality: 143613 bases at least Q30
Consensus quality: 148883 bases at least Q30
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Center code: BCM
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Birren, B., Linton, L., Nusbaum, C., Lander, E., Ali, A., Allen, N., Anderson, S., Barna, N., Bastien, V., Boguslavkiy, L., Boukhgalter, B., Anderson, S., Barna, N., Bastien, V., Boguslavkiy, L., Boukhgalter, B., Brown, A., Coak, C., Chazaro, B., Choepel, V., Colangelo, M., Collins, S., Collymore, A., Cook, A., Cooke, P., Dearellano, K., Dewar, K., Diaz, J.S., Dodge, S., Faro, S., Ferreira, P., Fitzhugh, W., Gage, D., Galagan, J., Gardyna, S., Ginde, S., Gord, S., Goyette, M., Graham, L., Grand-Pierre, N., Hagos, B., Horton, L., Hulme, W., Iliev, T., Johnson, R., Jones, C., Kamat, A., Karatas, A., Kells, C., LaRocque, K., Lamazares, R., Landers, T., Lehoczky, J., Levine, R., Liu, G., MacLean, C., Macdonald, P., Major, J., Marquis, N., Matthews, C., McCarthy, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Birren,B., Linton,L., Nusbaum,C. and Lander,E. Mus musculus, clone RP23-358A18
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On Aug 9, 2002 this sequence version replaced gi:20334333.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fujlyama, A., Yada, T., Totoki, Y., Watanabe, H. and Sakaki, Y. Homo sapiens genomic DNA Published Only in Database (2000) 2 (Dases 1 to 188172) 4 (Dases 1 to 188172) 5 (Dases 1 to 180172) 7 (Dases 1 to 180172) 8 (Dases 1 to 180172) 8 (Dases 1 to 180172) 9 (Dases 
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Fujiyama,A., Yada,T., Totoki,Y., Watanabe,H. and Sakaki,Y.
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COMMENT

This sequence will be replaced by the finished sequence as soon as it is available and

the accession number will be preserved

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* consists of 9 contigs. Gaps between the contigs
* are represented as runs of N. The order of the pieces
* is believed to be correct as given, however the sizes
* of the gaps between them are based on estimates that have
* provided by the submittor.
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                                                                                                                                                                                                                                * NOTE: This is a 'working draft' sequence. It currently
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (02-JUN-2002) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA On Jun 2, 2002 this sequence version replaced gi:19033670. All repeats were identified using RepeatMasker: Smit, A.F.A. & Green, P. (1996-1997) http://ftp.genome.washington.edu/RM/RepeatMasker.html
                                                                                                                                                                                                                                                                                                                    Sequencing vector: Plasmid; n/a; 100% of reads
Sequencing vector: Plasmid; n/a; 100% of reads
Chemistry: Dye-terminator Big Dye; 100% of reads
Assembly program: Phrap; version 0.960731
Consensus quality: 191646 bases at least Q40
Consensus quality: 193107 bases at least Q30
Consensus quality: 193517 bases at least Q20
Insert size: 194000; agarose-fp
Insert size: 193969; sum-of-contiss
Quality coverage: 6.0 in Q20 bases; agarose-fp
Quality coverage: 6.0 in Q20 bases; sum-of-contigs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Center project name: L23802
Center clone name: 358_A_18
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                                                                                                                                                                                                                                AC122259 209418 bp DNA linear HTG 23-1 Mus musculus chromosome UNK clone RP23-183E7, WORKING DRAFT SEQUENCE, 10 unordered pieces. AC122259 GI:21105101
                                                                                    1 (bases 1 to 209418) McPherson, J.D. and Waterston, R.H.
                                                                                                                         Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                      The sequence of Mus musculus clone Unpublished
2 (bases 1 to 209418)
McPherson, J.D. and Waterston, R.H.
                                                                                                                                                                                                                  HTG; HTGS_PHASE1; HTGS_DRAFT
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1981 2080: gap of 100 bp
2081 3501: contig of 1421 bp in length
3502 3601: gap of 100 bp
3602 468: contig of 3247 bp in length
6849 6948: gap of 100 bp
6949 11584: contig of 4636 bp in length
11585 11684: gap of 100 bp
11685 30031: contig of 18347 bp in length
30032 30131: gap of 100 bp
30132 88538: contig of 58407 bp in length
88539 88638: gap of 100 bp
88639 118825: contig of 30187 bp in length
118826 118925: contig of 30187 bp in length
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162364 162463: gap of 100 bp
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162464 . .194769
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/clone_lib="RPCI-23 Female
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequencing vector: M13; 08
Sequencing vector: plasmid; 1008
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Submitted (23-MAY-2002) Genome Sequencing Center, 4444 Forest Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Parkway, St. Louis, MO 63108, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Quality coverage: 7.49 in Q20 bases; agarose-fp Quality coverage: 6.37 in Q20 bases; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: submissions@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Web site:http://genome.wustl.edu/gsc/index.shtml
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Center code: WUGSC
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* runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       * NOTE: This is a 'working draft' sequence. It currently consists of 10 contigs. The true order of the pieces is not known and their order in this sequence record is
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125736. .192346
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88783. .125635
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                                                                    clone_end:SP6
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19848. .51248
                                                                                                                                                                                                                                                                                                           /clone="RP23-183E7"
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    /note="assembly_name:Contig15
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2: gap of unknown length
7: contig of 13165 bp in length
7: gap of unknown length
8: contig of 31401 bp in length
8: gap of unknown length
2: contig of 37334 bp in length
2: contig of 37334 bp in length
5: gap of unknown length
6: contig of 36853 bp in length
6: contig of 66611 bp in length
6: contig of 1046 bp in length
6: contig of 1046 bp in length
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6: gap of unknown length
6: contig of 4496 bp in length
8: contig of 4968 bp in length
8: contig of 4978 bp in length
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/note="assembly_name:Contig7"
198189..203515
/note="assembly_name:Contig8"
eature 203616..209418
/note="assembly_name:Contig9"
58209 a 45087 c 44929 g 60292 t
                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (23-OCT-2001) DOE Joint Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 94598, USA
On Oct 23, 2001 this sequence version replaced gi:7711412.
Draft Sequence Produced by DOE Joint Genome Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (15-SEP-1999) Production Sequencing Facility, DOE Joint Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 94598, USA 3 (bases 1 to 238472) DOE Joint Genome Institute and Stanford Human Genome Center.
                                                                                                                                                                                                                                                                                                                 Quality: Phrap Quality >=40 99.7% of Sequence; Estimated Total Number of Errors is 0.5.
                                                                                                                                                                                                                                                                                                                                                      www-shgc.stanford.edu
                                                                                                                                                                                                                                                                                                                                                                         Finishing Completed at Stanford Human Genome Center
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DOE Joint Genome Institute.
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1 (bases 1 to 238472)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished
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                                                                                                                                                                                                                /db_xref="taxon:9606"
/chromosome="5"
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DEFINITION
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Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T.,
Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T.,
Barbarla, J., Benton, J., Binage, K., Blankenburg, K., Bonnin, D.,
Bunck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P.,
Buck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P.,
Buck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P.,
Buck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P.,
Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C.,
Cleveland, C.D., Cox, C., Coyle, M., B., David, R.,
Davila, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A.,
Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H.,
Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Dutbin, K.J.,
Barnhart, C., Edgar, D., Edwards, C.C., Blaj, C., Escotto, M.,
Earnhart, C., Edgar, D., Edwards, C.C., Blaj, C., Escotto, M.,
Falls, T., Ferraguto, D., Flagg, M., Ford, J., Foster, P., Frantz, P.,
Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R.,
Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Hernandez, J.,
Hernandez, C., Hodgson, A., Hogues, M., Holloway, C., Hollins, B.,
Homsi, F., Howard, S., Huber, J., Hulyk, S., Hume, J., Jackson, L.E.,
Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S.,
Kally, S., Khan, U., King, L., Koverh, J., Kovar, C.,
Kratovic, J., Kureshi, A., Landry, N., Leal, B., Lewis, L.C., Lewis, L.,
Lid, J., Li, Z., Lichtarge, O., Lieu, J., Liu, W., Martinez, E.,
Mahesiwari, M., Mapua, P., Martin, R., Mardinde, A., Martinez, E.,
Mahesiwari, M., Mapua, P., Martin, R., Madon, M., Mei, G., Metzker, M.,
Mahser, G., Miner, Z., Mitchell, T., Mohabbat, K., Moygan, M., Morris, S.,
Noser, M., Poliokan, I., Rolfe, M., Ruiz, S., Savery, G.,
Scherer, S., Scott, G., Shen, H., Shoshtari, N., Sisson, I.,
Soderrer, S., Wattern, R., Parke, A., Thomas, N., Thomas, S.,
Usmani, K., Vasguez, L., Vera, V., Villalon, D., Vinson, R., Wang, Q.,
Wall, S., Marten, R., Washington, C., Watlington, S
                                                                                                                        Submitted (11-JUL-2002) Human Genome Sequencing of Molecular and Human Genetics, Baylor College Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                     Worley,K.C.
Direct Submission
Submitted (17-SEP-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
3 (bases 1 to 239904)
                                                                                                                                                                                                                                                                                       Worley, K.C
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA linear HTG 11-JUL-2002 Rattus norvegicus clone CH230-162p21, *** SEQUENCING IN PROGRESS ***, 44 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rattus
                                                                             Jul 10, 2002 this sequence version replaced gi:18846093
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (bases 1 to 239904)
Center: Baylor College of Medicine
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                                          Genome Center
                                                                                                                                                          Center, Department of Medicine, One
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Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu

Center code: BCM

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NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 44 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.

This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Center project name: GEXO
Center clone name: CH230-162P21
Center clone name: CH230-162P21
Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of read Assembly program: Phrap; version 0.990329
Consensus quality: 19179 bases at least 040
Consensus quality: 196736 bases at least 030
Consensus quality: 201345 bases at least 020
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1 (bases 1 to 247865)

Muzny, D.M., Adams, C., Adio-Oduola, B., Ali-osman, F.R., Allen, Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T. Barbaria, J., Benton, J., Bimage, K., Blankenburg, K., Bonnin, D. Bouck, J., Bowle, S., Brieva, M., Brown, E., Brown, M., Bryant, N. Buhay, C., Burch, P., Burkett, C., Burrell, K.L., Byrd, N.C., Carron, T.F., Carter, M., Cavazos, S.R., Chaeko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C.,
                                                                                                                                                                                                                                                                                                                                                                        Similarity
19; Conser
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/db_xref="taxon:10116"
/clone="CH230-162P21"
49743 c 49983 g 64401 t
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                                                                                                                          Eutheria;
                                                                                                                                       Metazoa;
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170277: contig of 5772 bp in length
170377: gap of unknown length
179323: contig of 8946 bp in length
179423: gap of unknown length
193428: contig of 14075 bp in length
193598: gap of unknown length
208152: contig of 14554 bp in length
208152: gap of unknown length
208252: gap of unknown length
208252: gap of unknown length
208331: contig of 14879 bp in length
223331: gap of unknown length
233331: gap of unknown length
233904: contig of 16673 bp in length
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143648: contig o
143748: gap of u
154717: contig o
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81606: gap of
87909: contig
88009: gap of
94461: contig
94561: gap of
101192: contig
101292: gap of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        154817: gap of 164405: contig
                                                                                                                    Chordata; Craniata; Vertebrata; Euteleostomi; Rodentia; Sciurognathi; Muridae; Murinae;
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Cleveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Davila, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwalte, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Falls, T., Ferraguto, D., Elwards, C.C., Elhaj, C., Escotto, M., Falls, T., Ferraguto, D., Elwards, C.C., Elhaj, C., Escotto, M., Falls, T., Ferraguto, D., Elwards, C.C., Elhaj, C., Escotto, M., Frantz, P., Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Gorrell, J.H., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K., Harris, C., Hodgson, A., Hogues, M., Holloway, C., Halliton, K., Honsoi, F., Howard, S., Huber, J., Hulyk, S., Hume, J., Jackson, L.E., Jackson, E., Kally, S., Khan, U., King, L., Korvah, J., Kovar, C., Karlsson, E., Kelly, S., Khan, U., King, L., Korvah, J., Kovar, C., Kratovic, J., Kureshi, A., Landry, N., Leal, B., Lewis, L.C., Lewis, L., Li, W., Loulsey, A., Holloway, C., Martindale, A., Martindale, A., Martinda, R., Ma, J., Maheshwari, M., Mapua, P., Martin, R., Martindale, A., Martinda, E., Mansey, E., Mawhiney, E., McLeod, M.P., Martindale, A., Martinda, E., Martinda, R., Ma, J., Nguyen, N., Nickerson, E., Nwokenkwo, S., Oguh, M., Okwuonu, G., Oragunye, N., Ovledo, R., Pace, A., Payton, B., Peery, J., Perez, L., Pickens, R., Primus, E., Pu, L.L., Quiles, M., Ren, Y., Rheer, S., Sott, G., Shen, H., Shooshtari, N., Sisson, L., Sher, H., Shooshtari, N., Sisson, H., Shooshtari, N., Sisson, H., Shooshtari, N., Sisson, H., Shooshtari, N., Sisson, T., Wang, S., Ward-Moore, S., Warren, R., Waoden, S., Watliamas, G., Williamon, A., Wuczyk, R., Wooden, S., Watlingson, A., Whore, S., Own, S., Watlingson, A., Whore, S., Own, S., Watlingson, A., Waczyk, R., Wooden, S., Watlingson, S., Watl
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On Jul
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Submitted (24-AUG-2002) Human Genome Sequencing Center, Depai of Molecular and Human Genetics, Baylor College of Medicine, Baylor Plaza, Houston, TX 77030, USA
Baylor Plaza, Houston, TX 77030 replaced qi:17063105.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of Molecular and Human
Baylor Plaza, Houston,
3 (bases 1 to 247865)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (11-OCT-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Worley, K.C.
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Worley, K.C
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                       (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 40 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                        NOTE: Estimated insert size may differ from sequence length
as soon as it is available and the accession number will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jul 24, 2002 this sequence version replaced gi:17063105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (bases 1 to 247865)
                                                                                                                                                                                                                                                                                                                                                                                                                                  Center project name: TUUI
Center clone name: CH230-2G16
Center clone name: CH230-2G16
Center clone name: Statistics
Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 211853 bases at least Q40
Consensus quality: 215254 bases at least Q30
Consensus quality: 217501 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Center code: BCM Web site: http://www.hgsc.bcm.tmc.edu/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Center: Baylor College
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REFERENCE AUTHORS

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* 174363 188460: contig of 14098 bp in length
* 188461 188560: gap of unknown length
* 203564 203563: contig of 15003 bp in length
* 203564 203663: gap of unknown length
* 203564 221346: contig of 17683 bp in length
* 221347 221346: contig of 17683 bp in length
* 221347 221346: gap of unknown length
* 221447 247865: contig of 26419 bp in length
* 221447 247865: contig of 26419 bp in length
* 20147 247865: contig of 26419 bp in length
* 20147865: core 19: degree of 26419 bp in length
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* 20147865: core 19: degree of 26419 bp in length
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Scoring table:
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and is derived
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                                                                                                                                                                                                                             Score
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/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA2002.DAT:*
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                                                                   cDNA encoding huma
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Human ovarian carc
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Ovarian carcinoma
  Ovarian carcinoma
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RESULT 1 AAF95007 ID AAF9

ALIGNMENTS

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Drosophila melanog	ABL15972	23	8268	4.9	18	45
	ABL13144	23	6399	4.9	18	44
	AAH54174	22	4106	4.9	18	43
•	AAH54943	22	3633	4.9	18	42
盲	AAI59883	22	3556	4.9	18	41
S. epidermidis gen	AAH54960	22	3281	4.9	18	40
DNA encoding nove	AAS91414	23	2253	4.9	18	39
Murine G713 encod	AAA55968	21	1791	4.9	18	ა 8
Human ORFX OR	AAC74983	21	1633	4.9	18	37
Staphylococcus ep	ABN92620	24	1596	4.9	18	<u>ა</u>
Staphylococcus	ABN91962	24	1587		18	ဌ
DNA encoding nove	AAS83729	23	1542		18	34
S. epidermidis ope	AAH52496	22	1152	4.9	18	ယ
Human immune/haema	AAK66340	22	484		18	32
Human immune/haema	AAK66339	22	484		18	31
Corn tassel-derive	ABL75647	24	286		18	30
	ABL76335	24	256		18	29
Corn tassel-derive	ABL75913	24	237	4.9	18	28
3	ABK62081	24	2222	5.1	19	27
	ABK62080	24	2080	•	19	26
Human reproductive	AAL07456	22	1226	•	19	25
Human ovarian canc	ABL81273	24	430	•	55	24
Ovarian carcinoma	ABL40351	24	480	31.7	117	23
Human ovarian	ABT03283	24	480	•	117	22
Human ovarian anti	ABQ54231	24	1608	•	142	21
Primer specific to	AAF94044	22	591	•	221	20
Ovarian carcinoma	ABL40350	24	1010	-	271	19
Human ovarian carc	ABT03282	24	1010	•	271	18
Ovarian carcinoma	ABL48768	24	396	•	281	17
Human ovarian care	ABT03085	24	396	76.2	281	16
	AAF94818	22	396	•	281	15
Human secreted	AAF64188	22	1956	76.7	283	
Human secreted	AAF22400	21	1953	•	283	13
Human G protein co	AAD18690	22	2528	•	314	12
	ABL40352	24	1897	85.4	315	1
Human ovarian car	ABT03284	24	1897	85.4	315	0

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AAF95007 standard; DNA; 369 BP.

AAF95007;

23-MAY-2001 (first entry)

Human ovarian cancer associated coding sequence SEQ ID NO: 199

Human, ovarian cancer; vaccine; gene therapy; carcinoma; ds.

Homo sapiens.

W0200118046-A2.

15-MAR-2001.

08-SEP-2000; 2000WO-US24827.

10-SEP-1999; 99US-0394374.
01-MAY-2000; 2000US-05561778.
15-AUG-2000; 2000US-05561778.
15-AUG-2000; 2000US-0556668.

(CORI-) CORIXA CORP.

Isolated polypeptides associated with ovarian carcinomas, and the

WPI; 2001-211395/21.

Xu J,

Stolk JA;

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RESULT 2
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                                                                                                                                                                                      (first entry)
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99US-0394374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 369; DB 24; Length 369; 100.0%; Pred. No. 5.5e-187;
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                                                                                                                                                                                                                                         ВP
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                                                                                                                                                                                                                                                                                                                                                          Q
                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC The invention relates to ovarian tumour polynucleotides and polypeptides CC that may be utilised in cancer therapy, for example in a vaccine or CC gene therapy. Polypeptides and polynucleotides of the invention are CC useful for detecting a cancer in a patient, for stimulating and/or CC expanding T-cells specific for a tumour protein, and for inhibiting the CC development of a cancer in a patient. They are also useful for cancer in a patient and for treating a cancer in a patient had for determining the presence of a cancer in a patient. CC The isolated polynucleotides of the invention are useful for their CC ability to selectively form duplex molecules with complementary stretches CC of the entire desired gene or gene fragments, and for designing and CC preparing ribozyme molecules for inhibiting expression of tumour CC polypeptides in tumour cells. Polypeptides and polynucleotides of the CC invention are also useful in recombinant DNA molecules to direct CC expression of a polypeptide in appropriate host cells. The sequences CC overian carcinoma nroteins
                                                                                                                                                                                                                                                                                                                                                                                            Matches 369;
                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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15-AUG-2000;
07-SEP-2000;
14-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ovarian carcinoma proteins.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ovarian tumour polypeptide and prevention and/or treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ovarian tumour
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(STOL/)
(ALGA/)
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                             CACCATCCT
                                                                 CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCGCGCAC
                                                                                                      GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA
                                                                                                                                                                                                                                                                                        GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGGAGTTCATTGTAATTGCAC
                                                                                                                         CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA
                                                                                                                                                                                                                                                                        GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC
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 CACCATCCT
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                                                   CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCAC
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) ALGATE P A.
) FLING S P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                369 BP; 82 A; 109 C;
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                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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2000US-0640173.
2000US-0656668.
2000US-0713550.
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369
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                                                                                                                                                                                                                                                                                                                                                                                                                                               90 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polynucleotide useful in diagnosis, cancer, especially ovarian cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                86 T;
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ABK33543
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01-DEC-2000;
20-DEC-2000;
20-DEC-2000;
20-FEB-2001;
10-MAY-2001;
25-MAY-2001;
                                                                                                                                                                                                                                                                                            25-JUL-2000;
26-JUL-2000;
28-JUL-2000;
23-AUG-2000;
23-AUG-2000;
15-SEP-2000;
10-NGV-2000;
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25-JUL-2000;
25-JUL-2000;
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25-JUL-2000;
25-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; secreted protein; PRO; tubreast cancer; prostate tumour; pericyte cell proliferation; oth tumour necrosis factor alpha; gg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             te cell proliferation; chondrocyte cell proliferation; necrosis factor-alpha; gene; ss.
                                                                                                                                                                                                                                     2000US-220893P.
2000WO-US23710.
2000WO-US233522.
2000WO-US23328.
2000WO-US23328.
2000WO-US23873.
2000US-253646P.
2000US-2747259.
2000WO-US34956.
2001WG-US34956.
2001WG-US34956.
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2000US-220666P.
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2000US-220605P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRO; tumour; lung cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rectal tumour; liver
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    colon cancer;
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(GETH ) GENENTECH INC.

Baker KP, Grimaldi 'n, Desnoyers Gurney ΑĽ, Gerritsen Smith V, ME, Godd Stephan Goddard A, Watanabe Godowski PJ; CK, Wood WI;

P-PSDB; 2002-172001/22. AAU83599.

tumour or liver tumour -One hundred and twenty two nucleic acids encoding PRO polypeptides, useful for treating a PRO related disorder and for diagnosing tumours such as lung cancer, colon cancer, breast tumour, prostate tumour, rectal

Claim 2; Figure 15; 359pp; English.

The invention relates to one hundred and twenty two nucleic acids encoding PRO polypeptides. The sequences of the 122 PRO polynucleotides encode human secreted proteins. The PRO nucleic acids, polypeptides, agonists and antagonists are useful for treating a PRO related disorder. The PRO polypeptides are useful for diagnosing tumours, especially lung cancer, colon cancer, breast tumour, prostate tumour, rectal tumour or liver tumour. The PRO polypeptides are useful for stimulating the proliferation of, or gene expression, in pericyte cells, for stimulating the proliferation or differentiation of chondrocyte cells, for stimulating the release of tumour necrosis factor-alpha from human blood, for stimulating or inhibiting the proliferation of normal human dermal fibroblast cells. The PRO polypeptide may also be used as molecular weight markers and for tissue typing. The PRO nucleic acids have applications in molecular thallowed the proliferation of chords have applications in molecular biology, including use as hybridisation probes

RESULT 4

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RESULT 5
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03-APR-2001;
02-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1524
                                            Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                                                                   WPI;
                                                                                                     Xu J,
                                                                                                                                                                                                                                                       Human; ovarian cancer; cytostatic; gene; ss.
                                                                                                                                                                                                                                                                                    Human
                                                                                                                                                                                                                                                                                                      05-SEP-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                           ABT03277 standard; cDNA;
The present invention provides human ovarian
                   Claim
                                                                                                                                                                                13-NOV-2001; 2001WO-US45395
                                                                                                                                                                                                   23-MAY-2002
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les 315; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      in chromosome and gene mapping. \mathtt{ABK33536\text{-}ABK33657} represent human protein coding sequences of the invention.
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                                                                                   2002-500186/53.
                                                                                                                                                                                                                                                                                    ovarian
                  2;
                                                                                                     Stolk JA, Algate
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                                                                                                                                          2000US-0713550.
2001US-0825294.
2001US-0970966.
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                                                                                                                                                                                                                                                                                    carcinoma associated
                   189-190;
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                  197pp;
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Pred. No.
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                   English
                                                                                                                                                                                                                                                                   carcinoma;
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  cancer
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  proteins
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RESULT 6
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Best Local Similarity
Matches 315; Conserv
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                                                                                                                                                                                                                                                             Human; ovarian cancer;
cytostatic; gene; ss.
                                                                                                                                                                                                                                                                                             Human ovarian carcinoma
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                                    Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                                                              WPI;
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03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
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                                                                                                                                                                                                                         WO200239885-A2
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                                                                                                  Algate
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                                                                                                                                                                                                                                                                        ovarian carcinoma; gene therapy; immunotherapy;
                                                                                                                                                                                                                                                                                            associated coding sequence SEQ ID NO: 211.
                                                                                                                                                                                                                                                                                                                                                         1619
                                                                                                   PA,
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Pred No. 4.16
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                                                                                                   Fling
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4.1e-158;
                                                                                                   Molesh
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Claim

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Page 195;

197pp; English

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RESULT 7
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IID ABL4
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01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
                                                                                                                                                                             (XUJJ/)
(STOL/)
(ALGA/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ovarian carcinoma sequence isolate 57887 extended cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-JUN-2002 (first entry)
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  Ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                03-APR-2001;
                                                    WPI; 2002-171027/22.
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                                                                                                 Stolk JA, Algate
                                                                                                                                                  XU J.
) STOLK J A.
) ALGATE P A.
) FLING S P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer;
     tumour polypeptide and polynucleotide useful in
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2000US-0640173.
2000US-0656668.
2000US-0713550.
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                              RESULT 8
ABL40349
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Best Local
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                                                                                                                                                                                 Ovarian carcinoma 0591S nucleotide sequence.
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                                                              US2002004491-A1
                                                                                                                                                Ovarian cancer;
                                10-JAN-2002.
                                                                                               Homo sapiens
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                                                                                                                                              cancer therapy; vaccine; gene therapy; tumour; cancer;
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Pred. No. 4.1e-15
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03-APR-2001; 2001US-0825294

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01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
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330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1619 BP; 399 A; 415 C; 432 G; 373 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention relates to ovarian tumour polynucleotides and polypeptides
                                                              GCATCAGCTGCTACACACCCCCTCTTGTAACGGGCCAAGGCCCAAGAAAAGGGGAAGTT
                                                                                                                                 GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT
                                                                                                                                                                                                                         AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCGTCATCAGCGGCCT
CTGCCTCGGCCCTCA
                                                                                                                                                   GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT
                                                                                                                                                                                                                                                                                                                                               CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT
                                                                                                 GCATCAGCTGCTGCAACACCCCTCTTTGTAACGGGCCAAGGCCCCAAGAAAAGGGGAAGTT
                                                                                                                                                                                                         AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                                                                                                                                                                                                           GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
                                                                                                                                                                                                                                                                                                            GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STOLK J A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0394374.
2000US-0561778.
2000US-0640173.
2000US-0656668.
2000US-0713550.
                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    S P.
                                                                                                                                                                                                                                                                                                                                                                                                                                      85.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 315; DB 24;
Pred. No. 4.1e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fling SP;
                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 1619;
                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                  269
                                                                                                                                                                      269
                                                                                                                                                                                                         209
                                                                                                                                                                                                                                                                           149
                                                                                                                                                                                                                                                                                                                                               89
                                                                                                                                                                                                                                                                                                                                                                                                                   0
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AAF93845 ID AAF9 XX

AAF93845 standard; cDNA; 1890 BP

04 pp 04

150 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT

RESULT 9

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Matches 315;
                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    diagnosis of diseases associated with inappropriate secretory protein/membrane protein expression. The nucleic acids and complementary sequences may also be used as DNA probes in diagnostic assays (e.g. polymerase chain reactions (PCR)) to detect and quantitate the presence of similar nucleic acid sequences in samples. They may also be used to study the expression and function of secretory proteins/membrane polypeptides and their role in metabolism. The polypeptides may be used as antigens in the production of antibodies against them and in assays to identify modulators (agonists and antagonists) of expression and activity. The antibodies and antagonists may also be used as diagnostic agents for detecting the presence of the polypeptides in samples (e.g. by enzyme linked immunosorbant assay (ELISA). Examples of diseases which may be treated include rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention relates to nucleic acid sequences AAF93744 - AAF93916 which encode human secretory or membrane proteins represented by AAB88317 - AAB88419. Included in the invention are primers AAF93917 - AAF94295 and AAF62232 - AAF62235 which are used to isolate the cDNA sequences of the invention. The invention also includes methods for the production of antibodies directed against the proteins, and cDNA sequences, which can be used in vaccines. The polynucleotide sequences can be used in gene therapy. The polynucleotide sequences and the proteins they encode may be used in the prevention, treatment and approteins they encode may be used in the prevention, treatment and
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1890 BP; 419 A; 528 C; 533 G; 410 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              arthritis and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acids encoding secretory proteins/membrane proteins, useful gene therapy or as candidate target molecules in drug development -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P-PSDB; AAB88418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-JUL-2000; 2000EP-0114090
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          383
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        rheumatoid arthritis; diabetes; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human cDNA encoding a membrane or secretory protein clone PSEC0181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-MAY-2001
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                                                                                                                                  323 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 382
                                                                                                                                                                                                     30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT
                                                                                                                                                                                                                                                                                                               Local Similarity
                                   GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAAAGGTTCAAGAACATGTGTCAGAAAG
GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001-093989/11.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID 203; 609pp + CD ROM; English.
                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; 99JP-0194179.
; 2000JP-0118775.
; 2000JP-0183766.
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                                                                                                                                                                                                                                                                                                     85.4%; Score 315; DB 22; 100.0%; Pred. No. 4.2e-15
                                                                                                                                                                                                                                                                            0
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                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein;
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                                                                                                                                                                                                                                                                                                                                        Length 1890;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   drug
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RESULT 1
ABT03284
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                                                                                     Matches
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Best Local
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03-APR-2001;
02-OCT-2001;
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                                                                                                                                                                                                                                     Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                                                                                                                                                                                                                                                                                                                                                                              13-NOV-2001; 2001WO-US45395
                                                                                                                                                                                                                                                                                                                                                                                                                    23-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200239885-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human ovarian carcinoma associated coding sequence SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-SEP-2002 (first entry)
                                                                                                                                                              The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                                                                                                                                                                                        č
                                                                                                                               Sequence 1897 BP; 435 A; 521 C; 532 G; 407 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          623
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             503
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                                                                                                                                                     of the invention.
                                                                                                                                                                                                                                                                                                                              (CORI-) CORIXA CORP
 369
                                          309
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                    90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCATCAGCGGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTGCCTCGGCCCTCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTGCCTCGGCCCTCA 344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GTCTCATCGCCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT
                                         GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
                                                                                                                                                                                                          2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ovarian cancer; ovarian carcinoma;
                                                                                                                                                                                                                                                                                                        Stolk JA, Algate PA,
                                                                                               Similarity
                                                                                                                                                                                                          Page 196; 197pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene;
                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                   2001US-0825294.
2001US-0970966.
                                                                                                                                                                                                                                                                                                                                                                          200005-0713550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          637
                                                                                               85.4%; Score 315; 100.0%; Pred. No.
                                                                                      0,
                                                                                                                                                                                                                                                                                                          Fling
                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                           SP,
                                                                                                 4.2e-158;
                                                                                                            DB 24;
                                                                                                                                                                                                                                                                                                           Molesh DA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene therapy; immunotherapy;
                                                                                                          Length 1897;
                                                                                       Indels
                                                                                       0;
                                                                                       Gaps
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RESULT 11
ABL40352
ID ABL40
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    CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABL40352;
The invention relates to ovarian tumour polynucleotides and polypeptides that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient. They are also useful for the invention are useful for their
                                                                                                                                                                                                                                                                                                                                                                                                            US2002004491-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ovarian cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ovarian carcinoma 01034C/0591S consensus nucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABL40352 standard; cDNA; 1897 BP
                                                                                                                                                                                                                                                                                                                                                               03-APR-2001; 2001US-0825294.
                                                                                                                                                                                                                                                                                                                                                                                      10-JAN-2002.
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                                                                                                                                                                                                                                               (STOL/)
                                                                                                                                                                                                                                                                                                                 01-MAY-2000;
15-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     429
                                                                                                                    Claim 1a; Page 127-128; 131pp; English.
                                                                                                                                         prevention
                                                                                                                                                      Ovarian tumour
                                                                                                                                                                            P-PSDB;
                                                                                                                                                                                         WPI;
                                                                                                                                                                                                             Xu J,
                                                                                                                                                                                                                                                                                           14-NOV-2000;
                                                                                                                                                                                                                                                                                                       07-SEP-2000;
                                                                                                                                                                                                                                                                                                                                         10-SEP-1999;
                                                                                                                                                                                                                                   (FLIN/)
                                                                                                                                                                                                                                                                    (XUJJ/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sapiens
                                                                                                                                                                                       2002-171027/22.
                                                                                                                                                                                                            Stolk JA, Algate
                                                                                                                                                                                                                                   STOLK J A.

ALGATE P A

FLING S P.
                                                                                                                                                                            ABB09417
                                                                                                                                                                                                                                                                      ă
                                                                                                                                          mour polypeptide and polynucleotide useful in diagnosis, and/or treatment of cancer, especially ovarian cancer
                                                                                                                                                                                                                                                                                           2000US-0561778.
2000US-0640173.
2000US-0656668.
2000US-0713550.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                           99US-0394374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer therapy; vaccine; gene therapy; tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers 260..685
                                                                                                                                                                                                                                                                                                                                                                                                                                   /product=
                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag=
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                                                                                                                                                                                                              PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                     "Ovarian carcinoma protein
                                                                                                                                                                                                              Fling
                                                                                                                                                                                                                SP;
                                                                                                                                                                                                                                                                                                                                                                                                                                       01034C/0591S"
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                                                                                                                                                                                                                                                           Human: cardiovascular; tumourigenic disorder; aberrant angiogenesis; gene therapy; aberrant vascularisation; artherosclerosis; ovarian cancer; ischaemia/reperfusion injury; hypertension; arterial inflammation; psoriasis; endothelial cell disorder; diabetic retinopathy; restenosis; myocardial infarction; Grave's disease; G protein coupled receptor; GPCR 4941; vasotropic; hypotensive; antiinflammatory; cytostatic; antidiabetic; antipsoriatic; leukaemia; ss.
              25-APR-2001; 2001WO-US13788
                                                01-NOV-2001.
                                                                                WO200181634-A2
                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAD18690 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                          18-DEC-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the ovarian carcinoma 01034C/0591S consensus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1897 BP; 435 A; 521 C; 532 G; 407 T; 2 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT
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                                                                                                                                                                                                                                                                                                                                                                                                      protein coupled receptor (GPCR) 4941 cDNA.
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                                                                                                             /product= "Human GPCR 4941"
/note= "The CDS is specifically claimed in
the specification"
                                                                                                                                                                                  Location/Qualifiers 42..1403
                                                                                                                                                                /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 623
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AAF22400/c
ID AAF224
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DT 26-MAR
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KW Human;
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                                                                                                                                                                                                                                                                                                   1977
                                                                                                                                                                                                                                                                                                                                                                     2037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to a method for identifying a nucleic acid molecule (G protein coupled receptor gene, GPCR 4941) associated with a cardiovascular or tumourigenic disorder. The method comprising contacting a sample containing a nucleic acid molecule with a hybridisation probe or amplification primers and detecting the presence. The invention is used in gene therapy. The method of the invention is used for identifying nucleic acids or polypeptides associated with a cardiovascular or tumourigenic disorder such as aberrant angiogenesis, aberrant vascularisation, artherosclerosis, or ovarian cancer, ischaemia/ reperfusion injury, hypertension, restenosis, arterial inflammation, endothelial cell disorders, diabetic retinopathy, psoriasis, myocardial infarction, Grave's disease and leukaemia. The methods can also detect mRNA or genomic DNA in a sample. The present sequence is G protein
                                                                                                        AAF22400
                                                                                                                                       AAF22400 standard; cDNA;
                                                                                                                                                                                                                              1917 TGCCTCGGCCCTCA 1904
                                                                                                                                                                                                                                                                                                                                                                                                                                           2097
                                                                       26-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2217 AGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTG
                                                                                                                                                                                                                                                                331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2528 BP; 516 A; 766 C; 677 G; 567 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein coupled receptor (GPCR)-4941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Identifying nucleic acids for the diagnosis and treatment of cardiovascular and tumorigenic disorders, comprises identifying
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-APR-2000;
09-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                 211 TCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       151 AGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            91 CTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAGA 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31 AGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                          CATCAGCTGCTGCAACACCCCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGAAGTTC
                                                                                                                                                                                                                                                                TGCCTCGGCCCTCA 344
                                                                                                                                                                                                                                                                                                                                                                  TCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAGA
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                                    secreted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   314;
                                                                     (first entry)
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2000US-0635521.
                                  protein gene 28 SEQ ID NO:38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                85.1%;
                                                                                                                                           1953
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 314; DB 22;
Pred. No. 1.4e-15
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Human; secreted protein; diagnosis; immunosuppressive; antiarthritic;

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                                                                                                                                                                                                                                                                                              CC and polypeptides homologous to them. Human secreted protections have activities based on the tissues and cells the genes are expressed in. CC Examples of activities include: immunosuppressive; antiarthritic; CC antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; ccrebroprotective; nootropic; neuroprotective; antibacterial; virucide; CC fungicide; and ophthalmological. The polypucleotides and proteins can be used to prevent, treat or ameliorate a medical condition in e.g. humans, CC mice, rabbits, goats, horses, cats, dogs, chickens or sheep. They are also used in diagnosing a pathological condition or susceptibility to a pathological condition. Disorders which are diagnosed or treated include also disorders e.g. neoplasms of the breast or liver, cardiovascular CC disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. corneal infection. The polypeptides can also be used to ald wound healing and epithelial cell proliferation, to prevent skin candiovascular conductaris. The polypeptides can also be used to chamber the summan of the interest of transplantation, for comportant conducture of primary lissues, to regenerate tissues and in chamber to increase or decrease storage capabilities. AAF22364 to the present invention are researt sequences used in the exemplification of the present invention.
                                                                                                                                                                 Matches
                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; ophthalmological; gene therapy; autoimmune disease; neoplasm; rheumatoid arthritis; hyperproliferative disorder; cardiac arrest; cardiovascular disorder; cerebrovascular disorder; cerebrovascular disorder; cerebrovascular disorder; disease; infection; angiogenesis; nervous system disorder; Alzheimer's disease; infection; ocular disorder; corneal infection; wound healing; skin aging;
  1585
                                                                                1645
                                                                                                                                                                                                                                           Sequence 1953 BP; 428 A; 540 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF22373 to AAF22421 encode the human secreted proteins given in AAB6313 to AAB63183. AAB63183 to AAB63231 represent more human secreted proteins
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolated nucleic acid molecule encoding a human secreted protein used in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-APR-1999;
20-JAN-2000;
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                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                     present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                additive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2000-647420/62.
                   GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
                                                                                                                    CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 89
GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
                                                                            CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 1586
                                                                                                                                                                 283;
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                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page 440; 533pp; English.
                                                                                                                                                                 Conservative
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2000US-0176931.
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                                                                                                                                                                                 76.7%;
100.0%;
                                                                                                                                                              0;
                                                                                                                                                                               Score 283;
Pred. No.
                                                                                                                                                                                                                                           524 G; 458 T; 3 other;
                                                                                                                                                              Mismatches
                                                                                                                                                                                                       DB 21;
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             CC Human secreted proteins AAB75506 - AAB75554 are encoded by polynucleotide CC sequences AAF64176 - AAF64224. The specification includes amino acid CC sequences AAB75555 - AAB75606 which represent fragments of the human CC secreted proteins, and protein sequences with which they share homology. CC The proteins and polynucleotides, their agonists and antagonists have CC activities dependent on the tissues and cells in which they are CC expressed, examples of these activities include, immunosuppressive; CC unitarthritic; antirheumatic; antiproliferative; cytostatic; cardiant; CC virucide; fungicide; opthalmalogical; and vulnerary. The proteins, CC virucide; fungicide; opthalmalogical; and vulnerary. The proteins of the polynucleotides, agonists and antagonists can be used to treat or diagnose various diseases and disorders including, autoimmune CC diseases e.g. rheumatoid arthritis, hyperproliferative disorders CC e.g. neoplasms of the breast or liver, cardiovascular disorders (e.g. cardiac arrest, cerebrovascular disorders e.g. Alzheimer's disease, functions caused by bacteria, viruses and fungi and ocular disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; opthalmalogical; vulnerary; autoimmune disease; cardiovascular disorder hyperproliferative disorders; cerebrovascular disorder; wound healing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1405
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                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 443-444; 542pp; English
                                                                                                                                                                                                                                                                                                                                                                                      preventing, treating or ameliorat
Parkinson's diseases and cancers
                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid molecules encoding human secreted proteins, used in preventing, treating or ameliorating a disorder, e.g. Alzheimer's and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rosen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ROSE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nervous system disorder; ocular disorder; skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; secreted protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted protein gene 13 SEQ ID NO:23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-APR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      additive; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GCATCAGCTGCTGCAACACCCCTCTTGTAACGGGCCAAGGCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB75518.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HUMAN GENOME SCI INC
ROSEN C A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ruben SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000WO-US14973
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0138630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Komatsoulis GA;
The polypeptides can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunosuppressive; antiarthritic; antirheumatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chemotaxis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disorder;
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e.g. corneal infection.

also

used

to aid wound

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RESULT 15
AAF94818
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Best Local S
Matches 283
                                                                                                                                                                 10-SEP-1999;
01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                healing and epithelial cell proliferation, to prevent skin ageing due to sunburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotraxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities. Included in the invention are polynucleotide sequences AAF64167 - AAF64175 and peptide AAF7505 which are used in the isolation, identification and characterisation of the proteins of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1405
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                                            Isolated polypeptides associated with ovarian carcinomas, nucleic acids that encode them, useful for the prevention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1525
            Claim 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF94818 standard;
                                                                                                                Хu J,
                                                                                                                                                                                                                                  08-SEP-2000; 2000WO-US24827.
                                                                                                                                                                                                                                                            15-MAR-2001
                                                                                                                                                                                                                                                                                      WO200118046-A2
                                                                                                                                                                                                                                                                                                                                        Human, ovarian
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                                                                                                                                                                                                                                                                                                                                                                                                                   AAF94818;
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                                                                                                                                         (CORI-) CORIXA CORP
                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     270
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                                                                                      2001-211395/21.
                                                                                                                Stolk JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
          Page 119;
                                   of ovarian cancers
                                                                                                                                                                ; 2000US-0561778.
; 2000US-0640173.
; 2000US-0656668.
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                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                              cancer associated coding sequence
                                                                                                                                                                                                                                                                                                                                        cancer;
                                                                                                                                                                                                          9908-0394374
         189pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA; 396
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                                                                                                                                                                                                                                                                                                                                        vaccine;
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Pred. No. 5.3e-141;
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                                                                                                                                                                                                                                                                                                                                       gene
                                                                                                                                                                                                                                                                                                                                      therapy;
                                                                                                                                                                                                                                                                                                                                        carcinoma; ss.
                                               carcinomas, and the prevention diagnosis
                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1363
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RESULT 16
ABT03085
IID ABT03
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ABT03085
IID ABT03
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AC ABT03
AC ABT03
AC Huma:
XX
AC Hum
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Best Local Simi
Matches 281;
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                                                                                                                                                           Novel ovarian cancer polypeptide and polynucleotide, useful fidetecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovar
                                                                                                                                                                                                                                                                                                                                                                                                                              14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
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                                                                                                                                                                                                                                                                                                                            Хu J,
                                                                                                                                                                                                                                                                                                                                                                             (CORI-) CORIXA CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ovarian carcinoma associated coding
                                                                                                                                                                                                                                                                                                                         Stolk JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                       Algate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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Pred. No.
                                                                                                                                                                                                                                                                                                                       Fling
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                                                                                                                                                                                                                                                                                                                            SP,
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6e-140;
                                                                                                                                                                                                                                                                                                                            Molesh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene therapy; immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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                                                                                                                                                                   ovarian
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                                                                                                                                                                      cancer
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Example 1; Page 116; 197pp; English.

present invention provides human over coding sequences. The sequences can

ovarian cancer can be used in t

the associated

sociated proteins diagnosis and

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RESULT 17
ABL48768
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Best Local :
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                                                                                                                                                                                     10-SEP-1999;
01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 396 BP; 90 A; 117 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                  (STOL/)
                                                                                                                                                                                                                                                                                                                                    Ovarian cancer; cancer therapy; vaccine; gene therapy; tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                         Ovarian carcinoma sequence isolate 21920
                                                                                                                                                                                                                                                                                                                                                                              18-JUN-2002 (first entry)
                                                  Ovarian tumour polypeptide and prevention and/or treatment of
                                                                                                                                                                                                                                                                                                                                                                                                  ABL48768
                                                                                                                                                                                                                                                                                                                                                                                                                      ABL48768 standard; cDNA;
The invention relates to ovarian tumour that may be utilised in cancer therapy,
                             Claim 1a; Page 41-42; 131pp; English
                                                                                 WPI; 2002-171027/22.
                                                                                                     Хu J,
                                                                                                                                                                                                                                          03-APR-2001; 2001US-0825294.
                                                                                                                                                                                                                                                             10-JAN-2002.
                                                                                                                                                                                                                                                                                 US2002004491-A1
                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                             14-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          280
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCATCAGCTGCTAACACCCCTCTTTGTAACGGGCCAAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT 269
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                                    Stolk JA,
                                                                                                                                 STOLK J A.
ALGATE P A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                          FLING S P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                           99US-0394374.
2000US-0561778.
2000US-0640173.
2000US-0656668.
2000US-0713550.
                                                                                                     Algate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         76.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                       396
                                                                                                     PΑ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 281;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95 G; 92 T; 2 other;
                                                                                                     Fling
                                                  polynucleotide useful in diagnosis, cancer, especially ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6e-140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 24;
polynucleotides and polypeptides for example in a vaccine or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 396;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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RESULT 18
ABT03282
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for a stimulating an immune response in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The sequences given in records ABL48760-ABL48956 represent polynucleotides encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 396 BP; 90 A; 117 C; 95 G; 92 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ovarian carcinoma proteins.
                                                                                                                                                                                                                                                                                                                                                                                                           ABT03282 standard; cDNA; 1010 BP
                                                                                   14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
                                                                                                                                              13-NOV-2001; 2001WO-US45395
                                                                                                                                                                                                          WO200239885-A2
                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                     cytostatic;
                                                                                                                                                                                                                                                                                   Human;
                                                                                                                                                                                                                                                                                                                                               05-SEP-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                             ABT03282;
                            Хu J,
                                                                                                                                                                              23-MAY-2002
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                                                        (CORI-) CORIXA CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30 CAGGCTTTGCGCCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT 269
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GCATCAGCTGCTAACACCCCTCTTTGTAACGGGCCAAGG 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         281;
                                                                                                                                                                                                                                                                                                               ovarian carcinoma associated coding sequence SEQ ID NO: 212.
                                                                                                                                                                                                                                                                                   ovarian cancer; ovarian carcinoma;
                            Stolk JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                   gene; ss
                              Algate
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100.0%; Pred. No.
                              PΑ,
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                              Fling
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                              SP,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 24;
                              Molesh
                                                                                                                                                                                                                                                                                   gene therapy;
                              DA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 396;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                      immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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WPI; 2002-500186/53

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RESULT 19
ABL40350
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Best Local
                                                                             01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
14-NOV-2000;
                        (STOL/)
(ALGA/)
(FLIN/)
   Хu J,
                                                                                                                                             03-APR-2001; 2001US-0825294
                                                                                                                                                                                           US2002004491-A1.
                                                                                                                                                                                                                                                                   Ovarian carcinoma BF345141 nucleotide sequence.
                                                                                                                          10-SEP-1999;
                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                             Ovarian cancer; cancer therapy; vaccine;
                                                                                                                                                                                                                                                                                                                ABL40350;
                                                                                                                                                                                                                                                                                                                                     ABL40350 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                        (XUJJ/)
                                                                                                                                                                     10-JAN-2002
                                                                                                                                                                                                                                                                                         28-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1010 BP; 266 A; 344 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 2; Page 195; 197pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                            498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 378
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31 AGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTG 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ocal Similarity
es 271; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                     CATCAGCTGCTGCAACACCCCTCTTTGTAAC 301
                                                                                                                                                                                                                                                                                                                                                                                                                                 CATCAGCTGCTACACCCCTCTTTGTAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTG
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                      ) STOLK J A.
) ALGATE P A.
) FLING S P.
  Stolk JA,
                                                        XU J.
                                                                            99US-0394374.
2000US-0561778.
2000US-0640173.
2000US-0656668.
2000US-0713550.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
 Algate
                                                                                                                                                                                                                                                                                                                                     cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             73.4%; Score 271; DI
100.0%; Pred. No. 1.
Live 0; Mismatches
  PA,
                                                                                                                                                                                                                                                                                                                                     1010
Fling
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                                                                                                                                                                                                                                            gene therapy; tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 1010;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         coding sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to ovarian tumour polynucleotides and polypeptides that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1010 BP; 266 A; 344 C; 254 G; 146 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ovarian tumour polypeptide and polynucleotide useful in diagnosis, prevention and/or treatment of cancer, especially ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim la; Page 126-127; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI;
      498
                                        271
                                                                               438
                                                                                                                   211
                                                                                                                                                            378
                                                                                                                                                                                                151
                                                                                                                                                                                                                                                                                                                258
                                                                                                                                                                                                                                                                                                                                31 AGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTG 90
                                                                                                                                                                                                                                                                         91
                                                                                                                                                                                                                                                                                                                                                                                                             Local
                      CATCAGCTGCTGCAACACCCCTCTTTGTAAC
                                                                                             TCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTG
                                                                                                                                                                                                                               AGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTG
                                                                                                                                                                                                                                                                     CTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAGA 150
                                                                                                                                                                                                                                                                                                          AGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTG 317
CATCAGCTGCTACACCCCCTCTTTGTAAC
                                                                         TCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTG
                                                                                                                                                    AGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2002-171027/22
                                                                                                                                                                                                                                                                                                                                                                                       271;
                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     73.4%; Score 271; DB 24
100.0%; Pred. No. 1.4e-1
                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
  528
                                      301
                                                                                                                                                                                                                                                                                                                                                                                                                       DB 24; Length 1010;
                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                               270
                                                                                                                                                                                        210
                                                                                                                                                    437
                                                                                                                                                                                                                                 377
                                                                           497
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0;

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RESULT 2
 Human; secretory protein; membrane protein; rheumatoid arthritis; diabetes; PCR primer;
                                                                                         Primer specific for DNA encoding secretory/membrane protein SEQ ID
07-JUL-2000;
                 10-JAN-2001
                                  EP1067182-A2
                                                                                                                                                AAF94044 standard;
                                                                                                              23-MAY-2001
                                                                                                                                                                 20
2000EP-0114090
                                                                                                             (first entry)
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vaccine; gene therapy;

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CC which encode human secretory or membrane protections represented by

CC AAR93917 - AAR94295 and AAR62232 - AAR62235 which are used to isolate the

CC AAR93917 - AAR94295 and AAR62232 - AAR62235 which are used to isolate the

CC CAAR93917 - AAR94295 and AAR62232 - AAR62235 which are used to isolate the

CC CAAR93917 - AAR94295 and AAR62232 which are used to isolate the

CC CDNA sequences of the invention. The invention also includes methods for

CC dequences, which can be used in vaccines. The polynucleotide sequences

CC can be used in gene therapy. The polynucleotide sequences and the

CC diagnosis of diseases associated with inappropriate secretory

CC proteins they encode may be used in the prevention, treatment and

CC diagnosis of diseases associated with inappropriate secretory

CC (e.g. polymerase chain reactions (CPCR)) to detect and quantitate the

CC (e.g. polymerase chain reactions (CPCR)) to detect and quantitate the

CC used to study the expression and function of secretory proteins/membrane

CC used to study the expression and function of secretory proteins/membrane

CC olympetides and their role in metabolism. The polympetides may be used

CC activity. The antibodies and antagonists against them and in assays to

CC dentify modulators (agonists and antagonists) of expression and

CC activity. The antibodies and antagonists may also be used as therapeutic

CC agents to down regulate expression and activity. The antibodies may also

be used as diagnostic agents for detecting the presence of the

CC polypeptides in samples (e.g. by enzyme linked immunosorbant assay

CC (ELISA). Examples of diseases which may be treated include rheumatoid

CC arthritis and diabetes.
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BXBXAXB
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                                                                                                                                     ABQ54231
                                                                                                                                                        RESULT 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 221;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; SEQ ID 478; 609pp + CD ROM; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acids encoding secretory proteins/membrane progene therapy or as candidate target molecules in drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-093989/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention relates to nucleic acid sequences AAF93744 - AAF93916
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 591 BP;
Human ovarian antigen HVCAG04 cDNA,
                                       22-AUG-2002
                                                                                                                   ABQ54231 standard; cDNA; 1608
                                                                                                                                                                                                                                                                                                                                                                                                                                                             323
                                                                                                                                                                                                                                                                                                                                                                              383
                                                                                                                                                                                                                     503
                                                                                                                                                                                                                                                          210
                                                                                                                                                                                                                                                                                                                                                                           90 GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT
                                                                                                                                                                                                                     GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCA 543
                                                                                                                                                                                                                                        GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCA 250
                                                                                                                                                                                                                                                                                                AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                                                                                                                                                                                                                                                                      AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Isogai T, Nishikawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HELIX RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; 99JP-0194179.
; 2000JP-0118775.
; 2000JP-0183766.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      108 A; 198 C; 173 G; 109 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                59.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 221; DB 22,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ţ,
                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kawai Y,
  SEQ ID NO:111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sugiyama T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 591;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             proteins, useful in
ruq development -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        development
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hayashi K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                     149
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                                                                                                                                                                                                                                                                                                                                                                              442
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CC ABP43228) and to cDNAs encoding them (ABQ54131-ABQ56305), and also CC encompasses polypeptides 90% identical and polynucleotides 95% identical contents on the sequences of the invention. The invention additionally relates to recombinant vectors and host cells comprising human ovarian antigen CC polynucleotides, antibodies against human ovarian antigen CC inventing, prognosing or preventing various ovary and/or breast-related CC metastaric tumours of ovarian or breast origin, reproductive system CC disorders. Such conditions include ovarian cancer and breast cancer, and CC metastaric tumours of ovarian or breast origin, reproductive system CC disorders (e.g., infertility, disorders of pregnancy, anovulation, CC polycystic ovary syndrome, ovarian cysts, and dysmenorrhoea), endocrine climunodeficiencies, autoimmune oophoritis, and acquired vaginitis), immune disorders (e.g., congenital and acquired vaginitis), immune disorders (e.g., congenital and acquired contents of prepared disorders (e.g., anaemia), cardiovascular disorders, cophoritis, systemic lupus erythematosus), CC invention of varian antigen expression or activity. The polynucleotides may also be used in screening for compounds which modulate ovarian antigen expression or activity. The polynucleotides may be used as food additives or to prepare antibodies useful in disease diagnosis, drug targeting and phenotyping. The present convention of individuals and in forensic analysis, and the convention of individuals and in forensic analysis, and the present sequence represents cDNA encoding a human ovarian antigen of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; ovarian antigen; ovary; ovarian; breast; cancer; tumour; ovarian cancer; breast cancer; tumour; reproductive system disorder; infertility; pregnancy disorder; anovulation; polycystic ovary syndrome; pcOS; ovarian cyst; dysmenorrhoea; endocrine disorder; infection; inflammatory condition; immune disorder; blood disorder; cardiovascular disorder; respiratory disorder; neurological disorder; gastrointestinal disorder; urinary system disorder; drug screening; gene therapy; chromosome mapping; forensic analysis; antibody preparation; cytostatic; immunomodulatory; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antibody preparation; cytostatic; immunomodulatory; neuroprotective; antilnflammatory; gynaecological; reproductive; chromosome 2q21-22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isolated nucleic acid molecules encoding novel ovarian polypeptides, useful in the prevention, treatment and diagnosis of cancer (e.g. ovarian cancer), immune disorders, cardiovascular disorders and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-JUN-2000; 2000US-209467P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-2001; 2001WO-US18569.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200200677-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID No 111; 2922pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   neurological diseases -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P-PSDB; ABP41154.
                                                                    Note: The sequence data for this patent did not form specification, but was obtained in electronic format
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2002-147878/19
                                                                      part of the printed directly from WIPO
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Query Match
Best Local Similarity
Matches 142; Conser

Conservative

38.5%; Score 142; D 100.0%; Pred. No. 1. tive 0; Mismatches

Sequence 1608

ftp.wipo.int/pub/published\_pct\_sequences.

BP; 402 A; 402 C; 417 G;

381 T; 6 DB 24; .2e-65;

other;

Length 1608;

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Gaps

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 22
                                                                                                                                         Matches 167;
                                                                                                                                                                 Query Match
                                                                                                                                                                                                                       The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                                                                                                                                                                                            Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian cancer \frac{1}{2}
                                                                                                                                                                                     Sequence 480 BP; 86 A; 154 C; 153 G; 87 T; 0 other;
                                                                                                                                                                                                                 of the invention.
                                                                                                                                                                                                                                                                           Claim 2; Page 196; 197pp; English.
                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-500186/53.
                                                                                                                                                                                                                                                                                                                                                                                          Хu J,
                                                                                                                                                                                                                                                                                                                                                                                                                                        14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-NOV-2001; 2001WO-US45395.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200239885-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytostatic; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; ovarian cancer; ovarian carcinoma; gene therapy; immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-SEP-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                 (CORI-) CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABT03283;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABT03283 standard;
150 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCAT
                                         369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         171
                                                                                         309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          291
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          231
                                                                90
                                                                                                             30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT
                                                                                                                                                      Local
                                     GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTGAGAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGATCATGTACCGCAAGTCCTGTGCATCAGCAGCCCTGTCTCATCACCCTCTGCCGGGT
                                                 GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
                                                                                       CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTCTTTGTAACGGCCAAGGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCTGCAACACCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTCTTTGTAACGGCCCAAGGCC 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ovarian carcinoma associated coding sequence SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                        Stolk JA, Algate PA,
                                                                                                                                                     Similarity
                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cDNA;
                                                                                                                                                    31.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      480
                                                                                                                                        0;
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                                                                                                                                                  Score 117;
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                       Fling SP,
                                                                                                                                       Mismatches
                                                                                                                                                  DB 24; Length 480; .8e-52;
                                                                                                                                                                                                                                                                                                                                                                                        Molesh DA;
                                                                                                                                       ۲.
                                                                                                                                     Indels
              197
                                                                                                                                       0;
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                                                                                                                                    Gaps
                                                                                    368
                                                                                                              89
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                                                                                                                                     0;
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Qy 밁

90

309

GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG

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Q
                                                                                                                     RESULT 23
ABL40351
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                                                                     Query Match
Best Local
                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (XUJJ/) XU J.
(STOL/) STOLK J A.
(ALGA/) ALGATE P A.
(FLIN/) FLING S P.
                                                                                                                                              useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the ovarian carcinoma BE336607 nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
14-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                         The invention relates to ovarian tumour polynucleotides and polypeptides that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are
                                                                                                                     Sequence 480 BP; 86 A; 154 C; 153 G; 87 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ovarian tumour polypeptide and polynucleotide useful in diagnosis, prevention and/or treatment of cancer, especially ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to ovarian tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim la; Page 127; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Xu J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-APR-2001; 2001US-0825294.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US2002004491-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ovarian cancer; cancer therapy; vaccine; gene therapy; tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ovarian carcinoma BE336607 nucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABL40351;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABL40351 standard; cDNA; 480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 429 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCAT 476
                   30
                                                                     Local Similarity
CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 89
                                                   167;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stolk JA, Algate PA,
                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2000US-0561778.
2000US-0640173.
2000US-0656668.
2000US-0713550.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99US-0394374
                                                                   31.7%;
                                                   0;
                                                                 Score 117; DB 24; Length 480; Pred. No. 2.8e-52;
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                                                   Mismatches
                                                 Indels
                                                 0,
                                                Gaps
                                                 0,
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RESULT 24
ABL81273
ID ABL81
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                                                                                                                                                                                                  The present invention describes a composition (I) comprising: carriers cc and immunostimulants; and a polypeptide (II) of a ovarian tumour cc polypeptide encoded by a polypucleotide (II) having a cDNA sequence cc (S1) from the 10912 nucleotide sequences as given in ABL/7923 to cc (S1) from the 10912 nucleotide sequences as given in ABL/7923 to cc (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to cc (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to cc (S1) can be used for detecting ovarian cancer in a patient's biological contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to patient, where the amount of polynucleotide hybridising to (IV) is cetected preferably by polymerase chain reaction (PCR). (I) comprising cc (III) and/or (II) is useful for stimulating and/or expanding T cells specific for an ovarian tumour protein comprising contacting T cells contacting for (IV) is useful in design and preparation of clls cuitable library e.g., a tumour cDNA library using well known
  В
                                    ş
                                                                                                Query Match
Best Local
                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABL81273;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABL81273 standard; cDNA; 430 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing polypeptide .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; ovarian cancer; ovarian tumour; cytostatic; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-MAY-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200192581-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human ovarian cancer related cDNA clone SEQ ID NO:4251.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID 4251; 489pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-122075/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Algate PA, Harlocker SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAY-2000; 2000US-207484P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-MAY-2001; 2001WO-US17756
                                                                                                                                                      Sequence 430 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              369 GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTGAGAAAG 428
AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCAT 476
                                                                              55.
                                                                                              Similarity
                                                                          14.9%;
ilarity 100.0%;
Conservative
                                                                                                                                                      92 A; 121 C; 108 G; 109 T; 0 other;
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                                                                            0;
                                                                                              Score 55;
Pred. No.
                                                                              Mismatches
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                                                                                              DB 24; 1
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                                                                                                              Length 430;
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                                                                            Gaps
                                                                              0;
  RESULT 25

RABIOTAT 5/5

RABIOTAT 5/5

RABIOTAT 5/5

RABIOTAT 5/5

RABIOTAT 10

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DT 21-NOV

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Human;

KW Cancer

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CANCER

XX

Human;

KW Cancer

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Homo s

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19-MAY-2000;
07-JUN-2000;
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04-FEB-2000;
24-FEB-2000;
02-MAR-2000;
                                                                                                                                                                        30-AUG-2000;
01-SEP-2000;
01-SEP-2000;
01-SEP-2000;
01-SEP-2000;
05-SEP-2000;
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14-AUG-2000;
14-AUG-2000;
                                                            08-SEP-2000;
08-SEP-2000;
08-SEP-2000;
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06-SEP-2000;
06-SEP-2000;
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14-AUG-2000;
18-AUG-2000;
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14-AUG-2000;
14-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; reproductive system related antigen; reproductive system disorder;
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22-AUG-2000;
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23-AUG-2000;
                                            08-SEP-2000;
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2000US-0217487.
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2000US-0216647
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2000US-0224519.
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2000US-0228924
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2000US-0225447.
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                                                                                                    2000US-0231242
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2000US-0230438
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PR 10-85EP-2000; 2000US-0233081.
PR 11-85EP-2000; 2000US-0233397
PR 14-85EP-2000; 2000US-0233399
PR 14-85EP-2000; 2000US-0233399
PR 14-85EP-2000; 2000US-0233404.
PR 14-85EP-2000; 2000US-0233409.
PR 14-85EP-2000; 2000US-0233404.
PR 14-85EP-2000; 2000US-0233409.
PR 14-85EP-2000; 2000US-0233404.
PR 12-85EP-2000; 2000US-0233404.
PR 25-85EP-2000; 2000US-023405.
PR 25-85EP-2000; 2000U
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RESULT 26
ABK62080/c
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Best Local :
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01-DEC-2000
05-DEC-2000
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05-DEC-2000
06-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
         11-AUG-2000;
19-OCT-2000;
                                                                                                                                                  Human; ss; gene; secreted protein; cancer; autoimmune disease; arthritis; osteoporosis; Alzheimer's disease; Parkinson's disease; meningitis; encephalitis; neoplasia; trauma; ischaemia; infarction; mania; stroke; cardiovascular disease; atherosclerosis; sepsis; anaemia; rheumatoid arthritis; hypothyroidism; allergic response; liver failure; multiple sclerosis; haemorrhage; paranola; obsessive compulsive disorder; autism; panic disorder; learning disability; feeding disorder; sleep pattern disorder; balance; perception; Thi-dependent insulitis; adult respiratory distress syndrome; ARDS.
                                                30-JUL-2001; 2001WO-US21124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention provides the protein and coding sequences of human reproductive system related antigens. These can tin the prevention and treatment of reproductive system disorders, including cancer. The present sequence is a genomic sequence enceprotein of the invention.
                                                                                                    WO200214358-A2
                                                                                                                           Homo sapiens.
                                                                          21-FEB-2002
                                                                                                                                                                                                                                                                               Human cDNA encoding novel secreted protein LP243(a).
                                                                                                                                                                                                                                                                                                                                    ABK62080;
                                                                                                                                                                                                                                                                                                                                                            ABK62080 standard;
                                                                                                                                                                                                                                                                                                         18-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1226 BP; 188 A; 438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated nucleic acid molecule encoding a reproductive system antigen is used in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-465570/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                       317 AAAAGGGGAAGTTCTGCCT 335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                         977 AAAAGGGGAAGTTCTGCCT 959
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           l Similarity
19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Barash SC,
          2000US-224642P
2000US-241779P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 10144; 1297pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-0251988
2000US-0256719
2000US-0251879
2000US-0251856
2000US-0251868
2000US-0251868
2000US-0251869
2000US-0251999
2000US-0254097
2000US-0254097
2001US-0259678
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2000US-0250391.
2000US-0251030.
                                                                                                                                                                                                                                                                                                                                                            cDNA;
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                                                                                                                                                                                                                                                                                                                                                            2080
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                            ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence encoding
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can be used
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ABK62081/c
ID ABK62081 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC The invention relates to a novel human secreted polypeptide having CC sequence 90% identical to the polypeptide sequences of LP105, LP061, CC LP224, LP230, LP239(a), LP243(a), LP243(b), LP253, LP218), LP251(a), CC LP224, LP230(b), LP233(a), LP255(a), LP244, LP186, LP251(b), LP255(b), CC or LP223(b), LP233(a), LP255(a), LP244, LP186, LP251(b), LP255(b), CC or LP223(b). Also included are the nucleic acids encoding the LP CC proteins (including complement, fragments encoding mature forms of the polypeptide or variant), a vector comprising the nucleic acid, a CC host cell comprising the vector, the preparation of the protein, CC an anti-LP antibody, ant/agonists of LP and anti-LP-encoding mRNA CC ribozymes. The secreted protein or its agonist is useful in the CC manufacture of a medicament for treating a mammal suffering from a CC disease (and in diagnosis), condition or disorder sasociated with CC aberrant levels of the secreted protein e.g. cancer, autoimmune disease, CC meningitis, encephalitis, neoplasia, trauma, ischaemia and infarction, CC mania, stroke, cardiovascular disease, atherosclerosis, rheumatoid CC sclerosis, hypothyroidism, anaemia, sepsis, allergic responses, multiple sclerosis, liver failure, haemorrhages, paranola, obsessive compulsive CC disorder, autism, panic disorder, learning disabilities, ALS (amylotrophic lateral sclerosis) psychoses, disorders in feeding, cspiratory distress syndrome (ARDS). The secreted protein is further CC useful for identifying compounds that bind to the secreted protein. The present sequence encodes a novel secreted protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                 Human; ss; gene; secreted protein; cancer; autoimmune disease; arthritis; osteoporosis; Alzheimer; s disease; Parkinson's disease; meningitis; encephalitis; neoplasia; trauma; ischaemia; infarction; mania; stroke; cardiovascular disease; atherosclerosis; sepsis; anaemia; rheumatoid arthritis; hypothyroidism; allergic response; liver failure; multiple sclerosis; haemorrhage; paranoia; obsessive compulsive disorder; autism; panic disorder; learning disability; feeding disorder; sleep pattern disorder; balance; perception; Th1-dependent insulitis;
                                    Homo sapiens
                                                                                                                                                                                                                                                                             Human cDNA encoding novel secreted protein LP243(b)
                                                                                                                                                                                                                                                                                                                            18-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1811 AAAAGGGGAAGTTCTGCCT 1793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2080 BP; 303 A; 758 C; 676 G; 342 T; 1 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel polypeptides and polynucleotides of secreted proteins useful for treating various diseases such as multiple sclerosis, cancer, autoimmune diseases, osteoporosis, Alzheimer's disease and Parkinson's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ELIL ) LILLY & CO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                        respiratory distress syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 Similarity
19; Conserv
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                                                                                                                                                                                                                                                                                                                       (first entry)
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Pred. No.
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14-MAY-2002 ABL75913;

(first

entry)

Corn; corn tassel-derived polynucleotide; cdps; hybrid breedinheritance; characteristic; growth; development; disease resenvironmental adaptability; quality; yield; molecular marker;

cdps; hybrid breeding; CDPs;

resistance;

trait; plant breeding; corn tassel; gene;

Corn tassel-derived polynucleotide (cdps) SEQ ID NO:5287

밁

1953

317 AAAAGGGGAAGTTCTGCCT 335

0

ABL75913 RESULT 28

ABL75913 standard; cDNA; 237

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The invention relates to a novel human secreted polypeptide having CC sequence 90% identical to the polypeptide sequences of LP105, LP061, CC LP224, LP240, LP239(a), LP243(a), LP243(b), LP253, LP218), LP251(a), CC LP252, LP239(b), LP223(a), LP255(a), LP244, LP186, LP251(b), LP255(b), CC LP252, LP239(b), LP223(a), LP255(a), LP244, LP186, LP251(b), LP255(b), CC proteins (including complement, fragments encoding mature forms of the CC proteins (including complement, fragments encoding mature forms of the CC proteins (including complement, fragments encoding mature forms of the protein, construction of the protein, construction of the protein, construction of the protein, cc an anti-LP antibody, ant/agonists of LP and anti-LP-encoding mana construction of the secreted protein or its agonist is useful in the construction of the secreted protein or disorder associated with cc antiference of a medicament for treating a mammal suffering from a construction or disorder associated with cc arthitis, osteoporosis, Alzheimer's disease, Parkinson's disease, carthitis, osteoporosis, Alzheimer's disease, Parkinson's disease, carthitis, stroke, cardiovascular disease, atherosclerosis, multiple contains, stroke, cardiovascular disease, atherosclerosis, multiple contents, bipothyroidism, anaemia, sepsis, allergic responses, multiple clarosis, panic disorder, learning disabilities, ALS (amylotrophic lateral sclerosis) psychoses, disorders in feeding, csepiratory distress syndrome (ARDS). The secreted protein is further useful for identifying compounds that bind to the secreted protein. The crespiratory construction and over secreted protein. The construction of the invention.
    Best Loc
Matches
                                            Query Match
                                                                              Sequence 2222 BP; 317 A; 827 C; 733 G; 345 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   autoimmune diseases, osteoporosis, Alzheimer's disease and Parkinson's
disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel polypeptides and polynucleotides of secreted proteins useful for treating various diseases such as multiple sclerosis, cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P-PSDB; AAU91329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-AUG-2000; 2000US-224642P.
19-OCT-2000; 2000US-241779P.
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                       Similarity
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      Conservative
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                     Score 19;
Pred. No.
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  Mismatches
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                   DB 24;
6.8;
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                                        Length 2222;
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               encode corn tassel-dérived polypeptides (CDPs). The cdps sequences (I) can be used for determining altered gene expression, to recover regulatory elements and to follow inheritance of desirable characteristics through hybrid breeding programs. (I) are also useful in the evaluation, and alteration of desired characteristics associated in the evaluation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence describes a purified corn tassel-derived polynucleotide sequence (cdps) comprising a nucleic acid sequenceted from those given in ABL70627 to ABL76833. The cdps selected from those given in ABL70627 to ABL76833.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel purified corn tassel-derived polynucleotide useful for determining altered gene expression, to recover regulatory elements and to follow inheritance of desirable characteristics through hybrid
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                                                Corn; corn tassel-derived polynucleotide; cdps; hybrid breeding; CDPs; inheritance; characteristic; growth; development; disease resistance; environmental adaptability; quality; yield; molecular marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 237
                                                                                                                       Corn tassel-derived polynucleotide (cdps) SEQ ID NO:5709.
                                                                                                                                                            14-MAY-2002
                                                                                                                                                                                                ABL76335;
                                                                                                                                                                                                                                   ABL76335 standard; cDNA; 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (or more) nucleic
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SHERMAN B K.
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                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 60 A; 64 C; 66 G; 46 T; 1 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequences.
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22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence describes a purified corn tassel-derived polynucleotide sequence (cdps) comprising a nucleic acid sequence selected from those given in ABL70627 to ABL70833. The cdps sequences encode corn tassel-derived polypeptides (CDPs). The cdps sequences (I) can be used for determining altered gene expression, to recover regulatory elements and to follow inheritance of desirable characteristics through hybrid breeding programs. (I) are also useful in the evaluation, and alteration of desirable characteristics through hybrid breeding programs.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel purified corn tassel-derived polynucleotide useful for determining altered gene expression, to recover regulatory elements to follow inheritance of desirable characteristics through hybrid
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(ITOL/) ITO L Y.
(SHER/) SHERMAN B K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-APR-1999;
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                                                inheritance; characteristic; growth; development; disease resistance; environmental adaptability; quality; yield; molecular marker; multigene trait; plant breeding; corn tassel; gene; ss.
                                                                                    Corn; corn tassel-derived polynucleotide; inheritance; characteristic; growth; devel
                                                                                                                                          Corn tassel-derived polynucleotide (cdps) SEQ ID NO:5021.
                                                                                                                                                                              14-MAY-2002
                                                                                                                                                                                                                 ABL75647
                                                                                                                                                                                                                                                   ABL75647 standard; cDNA; 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 256 BP; 72 A; 60 C;
                                                                                                                                                                                                                                                                                                                                                                               192 GTGCATCATCAGCGGCCT 209
                                                                                                                                                                                                                                                                                                                                           77
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18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          78 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          42 T; 4 other;
                                                                                                          cdps; hybrid breeding; CDPs;
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RESULT 31
AAK66339/c
ID AAK663
XX
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DT 06-NOV
XX
DE Human
E Human;
KW Human;
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Best Local S
Matches 18
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                           WO200157182-A2
                                                                          Human; immune; haematopoietic; immune/haematopoietic antigen; cancer; cytostatic; gene therapy; vaccine; metastasis: As
                                                                                                                 Human immune/haematopoietic antigen genomic sequence SEQ ID NO:21151
                                                                                                                                             06-NOV-2001
                                                                                                                                                                                              AAK66339 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             encode corn tassel-derived polypeptides (CDPs). The cdps sequences can be used for determining altered gene expression, to recover can be used for determining altered gene expression, to recover regulatory elements and to follow inheritance of desirable
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(SHER/) SHERMAN B K.
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                                                   Homo sapiens
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                   83 G;
                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                   52 T; 8 other;
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19-MAY-2000;
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2000US-0234997.
2000US-0234998.
2000US-0235484.
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2000US-0232399.
2000US-0232400.
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2000US-0231413
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                                                                  Rosen CA,
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                                                                  Barash SC,
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2000US -0236370
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2000US -0236611
2000US -023661
2000US -0236611
2000US -0236611
2000US -0236611
2000US -0236611

                                                                                                                                     GENOME
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RESULT 32
AAK66340/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 18
                                                         04-FEB-2000;

04-FEB-2000;

24-FEB-2000;

02-MAR-2000;

16-MAR-2000;

11-MAR-2000;

18-APR-2000;

19-MAY-2000;

19-MAY-2000;

107-JUN-2000;

28-JUN-2000;

30-JUN-2000;

30-JUL-2000;

11-JUL-2000;

11-JUL-2000;

11-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis -  \frac{1}{2} 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAK66340 standard; DNA; 484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human immune/haematopoietic antigen genomic sequence SEQ ID NO: 21152.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAK66340;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-JAN-2001; 2001WO-US01354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-NOV-2001
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2000US-0186350
2000US-0189874
2000US-0190076
2000US-0190123
2000US-0205515
2000US-0205467
2000US-0214886
2000US-0214887
2000US-0216847
2000US-0216847
2000US-0217487
2000US-0217487
2000US-0217487
2000US-02174863
2000US-02174863
2000US-02174863
2000US-02174867
2000US-02174863
2000US-02174863
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000US-0179065
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        therapy; vaccine; metastasis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.98;
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26-JUL-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000;

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08 - NOV - 2000;
09 - NOV - 2000;
17 - NOV - 2000;
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17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
01-DEC-2000
01-DEC-2000
05-DEC-2000
06-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
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17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
       AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the nucleic acids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotides may be used to prevent, and approach the munichaematopoietic-related diseases, especially axka103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-NOV-2000;
                                                                                                                                                                                                                                                       Nucleic acids encoding useful for preventing, metastasis -
                                                                                                                                                                                                                       Disclosure;
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2000US-0246523
2000US-0246524
2000US-0246525
2000US-0246526
2000US-0246527
2000US-0246609
2000US-0246611
2000US-0249209
2000US-0249209
2000US-0249209
2000US-0249210
2000US-0249210
2000US-0249211
2000US-0249211
2000US-0249212
2000US-0249213
2000US-0249216
2000US-0249264
2000US-0249264
2000US-0249264
2000US-02510391
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2000US-0246477
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                                                                                                                                                                                                                                                               human immune/hematopoietic diagnosing and/or treating
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 of
 haematopoietic-derived
                                                                                                                                                                                                                       Sequence Listing; English
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14-SEP-2000;
21-SEP-2000;
21-SEP-2000;
25-SEP-2000;
25-SEP-2000;
27-SEP-2000;
27-SEP-2000;
29-SEP-2000;
29-CCT-2000;
20-CCT-2000;
20-CC

2000US-0231243.
2000US-0231414.
2000US-0231413.
2000US-0232080.
2000US-0232080.
2000US-0232397.
2000US-0232401.
2000US-0232401.
2000US-0232401.
2000US-0232401.
2000US-0232401.
2000US-0234274.
2000US-0234274.
2000US-0234274.
2000US-0234299.
2000US-0234299.
2000US-0235834.
2000US-0235836.
2000US-0235836.
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2000US-0236368.
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2000US-0236369.
2000US-0237039.
2000US-0237039.
2000US-0237038.
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2000US-0237038.
2000US-0237038.
2000US-0237038.
2000US-0241785.
2000US-02441808.
2000US-0244617.
2000US-0244617.
2000US-0244617.

cancers

polypeptides, and

es, especially cells. AAK64703

14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
18-AUG-2000;
22-AUG-2000;
22-AUG-2000;
23-AUG-2000;
23-AUG-2000;
01-SEP-2000;

2000US-022964
2000US-0224519
2000US-0225213
2000US-0225214
2000US-0225267
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2000US-0225757
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Best Local Similarity
Matches 18; Conserv
                                       AAH55098 represent oligonucleotide sequences and primers which are used in the exemplification of the present invention.

N.B. The present invention specifically claims all the polynucleotide sequences given in the sequence listing of the present specification, however the sequence listing only goes up to SEQ ID NO:4454 so even though sequences are given in the disclosure for SEQ ID NO:4465 to 4472
                                                                                                                                                    containing them which are used to produce hosts cells which express the polypeptides. The polypeptides (II) (and/or nucleic acids) may then be used to vaccinate subjects and to raise antibodies against the bacteria. The polypeptides may also be used to assay for other inhibitors of their activity and therefore identify compounds that may be used for the treatment of S. epidermidis infections, e.g. endocarditis. AAH53971 to AAH55090 represent specifically claimed S. epidermidis genomic DNA polynucleotide sequences from the present invention. AAH55091 to
                                                                                                                                                                                                                                                                                                 AAH52304 to AAH53970 represent nucleic acids (I) encoding polyper (II), given in AAG81454 to AAG83120, from Staphylococcus epidermi (I) and (II) can have antibacterial activity and therefore can be in vaccination. The nucleic acids (I) may be used to produce the sepidermidis polypeptides (II) via the production of vectors
                              no sequences are present for SEQ ID NO:4455 to 4464.
                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 8; Page 144; 2188pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     S. epidermidis open reading frame nucleotide sequence SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kimmerly WJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GLAX ) GLAXO GROUP LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-NOV-2000; 2000WO-US30782.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        vaccination; endocarditis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Staphylococcus epidermidis SR1 strain; infection; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH52496
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 484 BP; 124 A; 110 C; 69 G; 181 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       represent sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      to AAK87694 represent human immune/haematopoietic antigen genomic sequences from the present invention. AAK54942 to AAK54950 and AAM82169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             457 CAGAAAGAAGTGATGGAG 440
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001-316495/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                       for vaccinating
                                                                                                                                                                                                                                                                                                                                                                                                                                              acids encoding polypeptides from Staphylococcus epidermidis, for vaccinating against infections, e.g. endocarditis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0164258
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 23;
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                                                                                                                                                                                                                              inhibitors of their
                                                                                                                                                                                                                                                                                                                                                            epidermidis
                                                                                                                                                                                                                                                                                                                                                                           polypeptides
                                                                                                                                                                                                                                                                                                                                         can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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Sequence 1152 BP;

429 A; 169 C;

241 G;

313 T; 0 other;

Best Local Similarity

100.0%;

Score 18; Pred. No.

DB 23; 23;

Length 1542;

4.98;

Query Match

Sequence 1542

BP;

416 A; 434 C; 433 G;

259 T; 0 other;

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The invention relates to isolated polynucleotide (I) and C polyneptide (II) sequences. (I) is useful as hybridisation probes, CC polynerase chain reaction (PCR) primers, oligomers, and for chromosome CC and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags CC for identifying expressed genes. (I) is useful in gene therapy techniques (C (II). (II) is useful for generating antibodies against it, detecting or compared activity of (II) or to treat disease states involving CC (II). (II) is useful for generating antibodies against it, detecting or compared to the compared to 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local
                          and to produce other types of data and products dependent on DNA and amino acid sequences. AAS64197-AAS94564 represent novel human diagnostic coding sequences of the invention.

Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutation responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          biodiversity
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23-AUG-2000; 2000US-0649167.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-FEB-2002 (first entry)
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nes 18; Conserv
ftp.wipo.int/pub/published_pct_sequences
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maging; diagnostic; genetic c
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Pred. No.
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08-NOV-1997;
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                                                                                                                                                                                                                                           antibacterial activity and can be used in gene therapy. The sequences can also be used in the diagnosis and treatment of bacterial infections, particularly S. epidermidis infections. The sequences can be used to screen for compounds able to interfere with the S. epidermidis life cycle or inhibit S. epidermidis infection.

N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-AUG-1998;
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                                                                                                                                                                                                                                                                                                                               ABN90538 to ABN93374 represent Staphylococcus epidermidis open reading frame (ORF) nucleic acid sequences which encode the amino acid sequences given in ABP35124 to ABP37960. The S. epidermidis sequences have
                                                                                                                                                                                                          Sequence 1587
                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID 1425;
                                                                                                                                                                                                                                                                                                                                                                                                       Novel isolated nucleic acid encoding a polypeptide, useful for diagnosing and
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 24-JUL-2002
                         ABN92620;
                                                  ABN92620
                                                                                                                                                                                                                                   USPTO web site.
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nes 18; Conserv
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                                                                                                             AATTCCAGCTGAACAACG 584
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                                                  standard;
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                                                                                                                                                                                                           325 G; 426 T; 0 other;
                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                        Staphylococcus epidermidis treating bacterial infections
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23;
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             Human; open reading frame; ORFX; detection; cytostatic; hepatotropic; vulnerary; antipsoriatic; antiparkinsonian; nootropic; neuroprotective; anticonvulsant; osteopathic; antiarthritic; immunosuppressant; cardiant; immunostimulant; thrombolytic; coagulant; vasotropic; antidiabetic; hypotensive; dermatological; immunosuppressive; antiinflammatory; antiviral; antibacterial; antifungal; antitheumatic; antithyoid; antianaemic; gene therapy; cancer; proliferative disorder; hypertension; neurodegenerative disease; diabetes mellitus; hypothyroidism; SCID; AIDS; coholesterol ester storage; systemic lupus erythematosus; infection; cholesterol ester storage; systemic lupus erythematosus; infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cycle or inhibit S. epidermidis infection.

N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel isolated nucleic acid encoding a Staphylococcus epidermidis polypeptide, useful for diagnosing and treating bacterial infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1596 BP; 616 A; 230 C; 327 G; 423 T; 0 other;
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   severe
                                                                                                                                                                                                Human ORFX ORF538 polynucleotide sequence SEQ ID NO:1075.
                                                                                                                                                                                                                                 08-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                  AAC74983 standard; cDNA; 1633
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
18; Conserv
combined immunodeficiency; malaria;
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gene therapy
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                                                                                                                                                                                                                                                                                                                                                                                                                                            100.08; F1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bush
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                267pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    open reading frame;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 18;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 24;
     autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORF; bacterial infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 1596;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO: 2083.
     disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
     asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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AAA55968/c
XEXTXA
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                                                                                                                                                                                                                                                                                                                                                                          RESULT 38
                                                                                                                                                                                                                                                                                                                                                                                                                     cc sequences have activities such as: cytostatic; hepatotropic; vulnerary; cc antipsoriatic; antiparkinsonian; nootropic; neuroprotective; osteopathic; anticonvulsant; antiarthritic; immunosuppressant; costopathic; anticonvulsant; thrombolytic; coagulant; vasotropic; coagulant; vasotropic; cantidiabetic; hypotensive; dermatological; immunosuppressive; antitinflammatory; antibacterial; antiviral; antifungal; antirheumatic; antithyroid; and antianaemic. The sequences can be used for determining the presence of or predisposition to, or preventing or treating contitions associated with an ORFX-associated disorder. The nucleic acids can be used to express ORFX proteins in gene therapy coroliferative disorders, neurodegenerative disorders, osteoarthritis, proliferative disorders, neurodegenerative disorders, osteoarthritis, cyafit vs host disease, cardiovascular disease, diabetes mellitus, hypertension, hypothyroidism, cholesterol ester storage, systemic lupus erythematosus, severe combined immunodeficiency (SCID), AIDS, viral, bacterial or fungal infection, malaria, autoimmune disorders, asthma, allergies, aplastic anaemia, burns, wounds, bone and cartilage damage, conductival haemoglobinuria, antinflammatory disease; to enhance
                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                   Matches
               Murine G713 encoding cDNA SEQ ID NO:6
                                                    05-SEP-2000
                                                                                          AAA55968;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAC74446 to AAC77606 encode the proteins given in AAB40237 to AAB43397, which represent the human ORFX open reading frames 1 to 3161. The ORFX
                                                                                                                             AAA55968 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                          Sequence 1633 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 5; Page 1013-1014; 5507pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel nucleic acids and peptides derived from open reading frame X, useful for treating e.g. cancers, proliferative disorders,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-602362/57.
P-PSDB; AAB40774.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-APR-1999; 99US-0127728.
30-MAR-2000; 2000US-0540763.
                                                                                                                                                                                                                                                                                                                                                                                                         coagulation; to inhibit thrombosis; and as a contraceptive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        neurodegenerative disorders and cardiovascular disease -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-MAR-1999;
02-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-MAR-2000; 2000WO-US08621.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           allergy; aplastic anaemia; nocturnal haemoglobinuria; burn; wound;
bone damage; cartilage damage; antiinflammatory disease; coagulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                         315 ACCAGTCCTTCTGCTCCC 332
                                                                                                                                                                                                                                                              231 ACCAGTCCTTCTGCTCCC 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           thrombosis; contraceptive; ss.
                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                 Conservative
                                                    (first entry)
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99US-0127636.
                                                                                                                                                                                                                                                                                                                                                                      335 A; 515 C; 480 G; 303 T; 0 other;
                                                                                                                                                                                                                                                                                        4.9%; Sur
100.0%; Pr
                                                                                                                                                                                                                                                                                                               Score 18; DB 21; Length 1633; Pred. No. 23;
                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                     RESULT
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                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                             Best
                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
  DNA encoding
                                                  13-FEB-2002
                                                                                                  AAS91414;
                                                                                                                                                  AAS91414 standard; cDNA; 2253 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1791 BP; 389 A; 531 C; 508 G; 363 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel polynucleotide of human G713 gene useful for diagnosis and prophylactic treatment of brain, psychiatric disorders like schizophrenia and bipolar disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; chromosome 13; G713; chromosome 13q31-q33; schizophrenia; biallelic marker; polymorphism; central nervous disease; detecti neuroleptic; G713 gene expression inhibitor; genotyping; brain disorder; psychiatric disorder; bipolar disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 234-236; 271pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-317979/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-OCT-1999;
                                                                                                                                                                                                                                                                         695
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-APR-2000
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                                                                                                                                                                                                                                                                                                                    84 ACGACTGCTCCTCCCCG 101
                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                    18;
                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                 Conservative
novel human diagnostic protein #27218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bougueleret L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98US-0103955
98US-0106457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99WO-IB01730
                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                       4.98;
                                                                                                                                                                                                                                                                         678
                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                               Score 18;
                                                                                                                                                                                                                                                                                                                                                                                           Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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23;
                                                                                                                                                                                                                                                                                                                                                                                                               21; Length 1791;
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EXXEXEX DX AX B
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AAH54960/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                        CC polypeptide (II) sequences. (I) is useful as hybridisation probes, CC polyperase chain reaction (PCR) primers, oligomers, and for chromosome cc and gene mapping, and in recombinant production of (II). The CC and gene mapping, and in recombinant production of (II). The CC and gene mapping, and in recombinant production of (II). The CC considers are also used in diagnostics as expressed sequence tags for identifying expressed genes. (I) is useful in gene therapy techniques cc for identifying expressed genes. (I) is useful in gene therapy techniques (II). (II) is useful for generating antibodies against it, detecting or CC quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical cimging of sites expressing (II). (I) and (II) are useful for treating CC imging of sites expressing (II). (I) and (II) are useful for treating CC the polypeptide and polynucleotide sequences have applications in CC diagnostics, forensics, gene mapping, identification of mutations cresponsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and CC amino acid sequences. AAS64197-AAS94564 represent novel human CC diagnostic coding sequences of the invention.

CC Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO at the commences.
                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local :
Staphylococcus epidermidis SR1 strain; infection; vaccination; endocarditis; {\tt ds.}
                                                          S
                                                                                            03-SEP-2001
                                                                                                                                 AAH54960
                                                                                                                                                               AAH54960 standard;
                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2253 BP; 541 A; 612 C; 527 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID No 27218; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to isolated polynucleotide (I) and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-MAR-2000;
23-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-MAR-2001; 2001WO-US08631
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; chromosome mapping; gene mapping; gene therapy; forensic; food supplement; medical imaging; diagnostic; genetic disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                        739
                                                                                                                                                                                                                                                                                          91 CTCCTCCCCGAGTTCAT 108
                                                        epidermidis
                                                                                                                                                                                                                                                                                                                                                                                                                                   ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                        CTCCTCCCCCGAGTTCAT 722
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2001-639362/73
                                                                                                                                                                                                                                                                                                                               18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RT,
                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2000US-0540217
2000US-0649167
                                                    genomic polynucleotide sequence SEQ ID NO:4324
                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tang
                                                                                                                                                                                                                                                                                                                                                                 4.98;
                                                                                                                                                                 3281
                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                          Score 18;
Pred. No.
                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                 573 T;
                                                                                                                                                                                                                                                                                                                                            DB
23;
                                                                                                                                                                                                                                                                                                                                                             23;
                                                                                                                                                                                                                                                                                                                                                                                                 0 other;
                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                             Length 2253;
                diagnosis;
                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                          0;
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RESULT 41
AAI59883
ID AAIS
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                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                              chemokinetic;
                                                                          peripheral nervous system; neuropathy; central nervous system; Calzheimer's; Parkinson's disease; Huntington's disease; haemosta amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic;
                                                                                                                     Human; nootropic; immunosuppressant; cytostatic; gene therapy; cancer;
                                                                                                                                                               Human polynucleotide SEQ ID NO 3872.
                                                                                                                                                                                                                                                                                                                                                  1118
                                                                                                                                                                                                                                 AAI59883;
                                                                                                                                                                                                                                                              AAI59883 standard; cDNA; 3556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 3281 BP; 1226 A; 464 C; 636 G;
                                              leukaemia; ss
                                                                                                                                                                                                22-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-316495/33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GLAX ) GLAXO
                                                                                                                                                                                                                                                                                                                                                                               269 TGCATCAGCTGCTGCAAC 286
                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                TGCATCAGCTGCTGCAAC 1101
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               acids encoding for vaccinating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 2062-2063; 2188pp; English.
                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                            thrombolytic; drug screening; arthritis; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GROUP
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                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                               4.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polypeptides from Staphylococcus epidermidis, against infections, e.g. endocarditis -
                                                                                                                                                                                                                                                                                                                                                                                                                0:
                                                                                                                                                                                                                                                                                                                                                                                                                                Score 18;
Pred. No.
                                                                                                                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                955 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                               DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                               22;
                                                                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 3281;
                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                          haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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RESULT 42
AAH54943/c
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                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                     Matches
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25-APR-2000;
09-JUL-2000;
19-JUL-2000;
03-AUG-2000;
                                                                                                                                                                                                                                                                                                      The invention relates to human nucleic acids (AAI57798-AAI61369) and the encoded polypeptides (AAM38642-AAM42213) with nootropic, immunosuppressant and cytostatic activity. The polynucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system diseases, such as Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activities such as: Immune system suppression, Activinyinhibin activity, chemotactic/chemokinetic activity, hammostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening, assays for receptor activity, arthritis and inflammation, leukaemias and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200153312-A1
                                                                                                                                                                                                                                                                                               assays for receptor activity, arthritis C.N.S disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; SEQ ID NO 3872; 10078pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel nucleic acids and such as central nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P-PSDB;
                                              Staphylococcus epidermidis SR1
                                                                 S. epidermidis
                                                                                      03-SEP-2001
                                                                                                          AAH54943;
                                                                                                                             AAH54943 standard; DNA; 3633
                                                                                                                                                                                                                                                         Sequence 3556 BP; 776 A; 1071 C; 1057 G; 652 T; 0 other;
                                                                                                                                                                                                                                                                            specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zhao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-SEP-2000;
19-OCT-2000;
WO200134809-A2
                  Staphylococcus epidermidis
                                      vaccination;
                                                                                                                                                                                                                                                                                      Note: The sequence data for this patent did not form part of the printed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HYSE-)
                                                                                                                                                                            629 ACCAGTCCTTCTGCTCCC 646
                                                                                                                                                                                                231 ACCAGTCCTTCTGCTCCC 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         , r A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2001-442253/47.
DB; AAM40727.
                                                                                                                                                                                                                    l Similarity
18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HYSEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Liu C,
Wang Z,
Zhou P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000US-0598042.
2000US-0620312.
2000US-0653450.
                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000US-0662191.
2000US-0693036.
2000US-0727344.
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2000US-0552317.
                                      endocarditis;
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                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INC
                                                                 genomic polynucleotide sequence SEQ ID NO:4307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Asundi V, Ch
Wehrman T, X
Goodrich R,
                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                       4.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               polypeptides, us
system injuries
                                       ds.
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xu C,
                                                                                                                                                                                                                    0
                                                                                                                                                                                                                             Score 18; pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nen R, Ma Y,
(u C, Xue AJ,
Drmanac RT;
                                                strain;
                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          useful for treating disorders
                                                infection; diagnosis;
                                                                                                                                                                                                                            DB
23;
                                                                                                                                                                                                                                      22;
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Yang Y,
                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                     Length 3556;
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Zhang
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                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                              RESULT 43
AAH54174/c
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Matches
                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-MAY-2001
           09-NOV-1999;
                                                 17-MAY-2001
                                                                                       Staphylococcus

 S. epidermidis

                                                                                                                                                           03-SEP-2001
           99US-0164258
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CC (II), given in AA631454 to AA63120, from Staphylococcus epidermidis. (I) and (II) can have antibacterial activity and therefore can be used CC (I) and (II) can have antibacterial activity and therefore can be used CC in vaccination. The nucleic acids (I) may be used to produce the CC s. epidermidis polypeptides (II) via the production of vectors containing them which are used to produce hosts cells which express the CC containing them which are used to produce hosts cells which express the CC used to vaccinate subjects and to raise antibodies against the bacteria. The polypeptides may also be used to assay for other inhibitors of their CC treatment of S. epidermidis infections, e.g. endocarditis. AAH53971 to CC AAH55090 represent specifically claimed S. epidermidis genomic DNA CC AAH55098 represent oligonucleotide sequences and primers which are used CC in the exemplification of the present invention. AAH55091 to AAH55098 represent invention specifically claims all the polynucleotide sequences given in the sequence listing of the present specification. CC consequences given in the sequence listing of the present specification, though sequences are present for SEQ ID NO:4454 so even consequences are present for SEQ ID NO:4455 to 4472, consequences are present for SEQ ID NO:4455 to 4461.
                                                                                                                                                                                                                                            Staphylococcus epidermias vaccination; endocarditis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAH54174 standard; DNA; 4106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1763 TGCATCAGCTGCTGCAAC 1746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 3633 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 8; Page 2042-2043; 2188pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acids encoding polypeptides from Staphylococcus epidermidis, useful for vaccinating against infections, e.g. endocarditis -
09-NOV-2000; 2000WO-US30782
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GLAX ) GLAXO GROUP LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               269 TGCATCAGCTGCTGCAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001-316495/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       l Similarity
18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000WO-US30782
                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                        epidermidis
                                                                                                                                                                                                                                                                                epidermidis
                                                                                                                                                                                                                                                                                                                                                 genomic polynucleotide sequence SEQ ID NO:3538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1340 A; 498 C; 707 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
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                                                                                                                                                                                                                                                         SR1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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Pred. No.
                                                                                                                                                                                                                                                                                      strain; infection; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB
23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1088 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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ABL13144/c ID ABL13144 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH52304 to AAH53970 represent nucleic acids (I) encoding polypeptides (II), given in AAG81454 to AAG83120, from Staphylococcus epidermidis. (C) and (II) can have antibacterial activity and therefore can be used to reactivation. The nucleic acids (I) may be used to produce the CC sepidermidis polypeptides (II) via the production of vectors (C) containing them which are used to produce hosts cells which express the CC used to vaccinate subjects and to raise antibodies against the bacteria. The polypeptides (II) and/or nucleic acids) may then be contained the production of vectors of their creatment of s. epidermidis infections, e.g. endocarditis. AAH53971 to AAH55090 represent specifically claimed s. epidermidis genomic DNA CC AAH55090 represent oligonucleotide sequences and primers which are used in the exemplification of the present invention. AAH55991 to CC AAH55098 represent oligonucleotide sequences and primers which are used in the exemplification of the present invention. CC N.B. The present invention specifically claimed s. phdermidis against the used considered sequences given in the sequence listing of the present specification, thouse the sequence listing of the present specification, though sequences are given in the disclosure for SEQ ID NO:4464 so even the sequence are present for SEQ ID NO:4465 to 4472, and the sequence are present for SEQ ID NO:4465 to 4472, and the sequence are present for SEQ ID NO:4465 to 4472, and the sequence are present for SEQ ID NO:4465 to 4472, and the sequence are present for SEQ ID NO:4465 to 4472, and the sequence are present for SEQ ID NO:4465 to 4472, and the sequence are present for SEQ ID NO:4465 to 4472, and the sequence are present for SEQ ID NO:4465 to 4472, and the sequence are present for SEQ ID NO:4465 to 4472, and the sequence are present for SEQ ID NO:4465 to 4472, and the sequence are present for SEQ ID NO:4465 to 4472, and the sequence are present for SEQ ID NO:4465 to 4472, and the sequence are present for SEQ ID NO:4465 to 4472, and the sequence are pr
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Best Local
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WPI; 2001-656860/75
P-PSDB; ABB69041.
                                                                                                     (PEKE ) PE
                                                                                                                                              23-MAR-2000;
11-JUL-2000;
                                                                                                                                                                                                         23-MAR-2001;
                                                                                                                                                                                                                                                  27-SEP-2001
                                                                                                                                                                                                                                                                                           WO200171042-A2
                                                                                                                                                                                                                                                                                                                                Drosophila melanogaster
                                                                                                                                                                                                                                                                                                                                                                       pharmaceutical; gene;
                                                                                                                                                                                                                                                                                                                                                                                  Drosophila; developmental biology; cell signalling; insecticide
                                                                                                                                                                                                                                                                                                                                                                                                                               Drosophila melanogaster expressed polynucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4106 BP; 1436 A; 659 C; 643 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            no sequences are present for SEQ ID NO:4455 to 4464.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acids encoding polypeptides from Staphylococcus epidermidis, useful for vaccinating against infections, e.g. endocarditis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        269 TGCATCAGCTGCTGCAAC 286
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                     CORP NY
                                                          Adams M,
                                                                                                                                                                                                         2001WO-US09231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                            2000US-191637P
2000US-0614150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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Pred. No.
                                                          Myers
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                                                          EW;
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Matches
                                                                     Query Match
Best Local :
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                                                                                                                                                                                                                                                insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABL16176-ABL30511), expressed DNA sequences (ABL01840-ABL16175) and the encoded proteins (ABB57737-ABB72072).
                                                                                                                                 Sequence 6399 BP; 1706 A; 1543 C; 1399 G; 1751 T; 0 other;
                                                                                                                                                                               at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                  The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 33914; 21pp + Sequence Listing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            interactions •
110 GTGAATTGCACGGTGAAC 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated nucleic acid
                                                18;
                                                                     Similarity
                                             Conservative
                                                                100.0%;
                                                                                           4.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          detection reagent for detecting 1000 or more for elucidating cell signalling and cell-cell
                                             0,
                                                                Score 18; Pred. No.
                                           Mismatches
                                                              DB
24;
                                                                                    23;
                                           0;
                                                                                    Length 6399;
                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                       English
                                                                                                                                                                                                                                                                                                                                                                                        The invention
                                        0,
                                        Gaps
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RESULT 45 ABL15972/c 5810 ABL15972 standard; 2 GTGAATTGCACGGTGAAC 5793 CDNA; 8268 ВP

В δ

Drosophila melanogaster Drosophila; developmental biology; cell signalling; insecticide; Drosophila melanogaster expressed polynucleotide SEQ ID NO 42398 pharmaceutical; gene; 26-MAR-2002 ABL15972; (first entry) ss.

W0200171042-A2 27-SEP-2001

23-MAR-2000; 11-JUL-2000; 2000US-191637P 2000US-0614150

23-MAR-2001; 2001WO-US09231

(PEKE ) PE CORP NY.

Venter JC, Adams M, Ŀ PWD, Myers EW;

2001-656860/75. ABB71869

New isolated nucleic acid genes from Drosophila and genes from Drosophila interactions detection reagent for detecting 1000 or more for elucidating cell signalling and cell-cell

Claim 1; SEQ ID NO 42398; 21pp + Sequence Listing; English

capable of detecting 1000 or more genes from Drosophila. The invention useful in developmental biology and in elucidating cell signalling and cell-cell interactions in higher eukaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences The invention relates to an isolated nucleic acid detection reagent capable of detecting 1000 or more genes from Drosophila. The invent (ABL16176-ABL30511), is

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Minimum DB
Maximum DB
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Result
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Word size
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               seq length: 0
seq length: 2000000000
      Issued_Patents_NA:*

1: /cgn2_6/ptodata/1

2: /cgn2_6/ptodata/1

3: /cgn2_6/ptodata/1

4: /cgn2_6/ptodata/1

5: /cgn2_6/ptodata/1

6: /cgn2_6/ptodata/1
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Match Length DB
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Copyright (c) 1993 - 2002 Compugen
        /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
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/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/backfiles1.seq:*
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US-09-134-001C-1425
US-09-134-001C-2083
US-09-134-001C-263
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US-08-532-8288-12
US-08-532-8288-1
US-08-532-8288-7
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US-08-967-104-7
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8184.096 Million cell updates/sec
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Sequence 1425, Apsequence 263, App sequence 263, App sequence 17, Appli sequence 12, Appli sequence 1, Appli sequence 1, Appli sequence 7, Appli sequence 9, Appli sequence 10, Appli sequence 10, Appli sequence 10, Appli sequence 11, Appli sequence 11, Appli sequence 21, Appli sequence 5, Appli sequence 5, Appli sequence 7, Appli sequence 17, Appli sequence 5, Appli sequence 5, Appli sequence 5, Appli sequence 5, Appli sequence 6, Appli sequence 6, Appli sequence 12, Appli sequence 13, Appli sequence 13, Appli sequence 13, Appli sequence 15, Appli sequence 15
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Sequence 2083, Application US/09134001C

Patent No. 6380370

GENERAL INFORMATION:

APPLICANT: Lynn Doucette-Stamm et al

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO S'

TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: GTC-007

CURRENT APPLICATION NUMBER: US/09/134,001C

CURRENT APPLICATION NUMBER: US 60/064,964

PRIOR APPLICATION NUMBER: US 60/055,779

PRIOR APPLICATION NUMBER: US 60/055,779

PRIOR APPLICATION NUMBER: US 60/055,779

PRIOR FILING DATE: 1997-08-14

SEQ ID NO 2083

LENGTH: 1596

TYPE: DNA

TO STAPHYLOCOCC

RESULT 2 US-09-134-001C-2083/c

37 10 10	Sequent Search S	RESULT	CC CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC
-09-134-001C-1425  Query Match Best Local Similarity 10 Matches 18; Conservativ 69 AATTCCAGCTGAACAACG	RESULT 1  VSequence 1425, Application US/0913, Patent No. 6380370  GENERAL INFORMATION: APPLICANT: Lynn Doucette-Stamm et TITLE OF INVENTION: NUCLEIC ACID. TITLE OF INVENTION: EPIDERMIDIS FILE REFERENCE: GTC-007  FILE REFERENCE: GTC-007  CURRENT APPLICATION NUMBER: US/09  CURRENT FILING DATE: 1998-08-13  PRIOR APPLICATION NUMBER: US 60/0  PRIOR FILING DATE: 1997-11-08  PRIOR APPLICATION NUMBER: US 60/0  PRIOR FILING DATE: 1997-08-14  NUMBER OF SEQ ID NOS: 5674  SEG ID NO 1425  LENGTH: 1587  TYPE: DNA  ORGANLESM: Staphylococcus epiderm  US-09-134-001C-1425	> /	17 17 17 16 16 16 16 16 16 16 17
tch al Similarity 10 18; Conservativ AAATTCCAGCTGAACAACG	Applicat ANTION: ANTION: ANTION: Van Doucet ENTION: CATION NO DE: GTC-00 DATE: NITON NUMB DATE: 199 ANTION NO ANTION: 199 DATE: 199	N N N N N N N N N N N N N N N N N N N	444444444444444444444444444444444444444
ANT: LYNN DOUGETTE-Stamm e OF INVENTION: MUCLEIC ACID OF INVENTION: MUCLEIC ACID OF INVENTION: MUCLEIC ACID OF INVENTION: MUCLEIC ACID OF INVENTION: MUCLEIC STORM IT FILING DATE: 1998-08-13 APPLICATION NUMBER: US 60/ FILING DATE: 1997-11-08 APPLICATION NUMBER: US 60/ FILING DATE: 1997-08-14 OF SEQ ID NOS: 5674 NO 1425 H: 1587 DNA APPLICATION S: 5674 INSM: Staphylococcus epideri- 001C-1425 Seal Similarity 100.0%; 18; Conservative 0; AATTCCAGCTGAACG 86	ation U ette-St NUCLEIC EPIDER 007 007 1198- 1198- 1298- 1297-08- 15674		1643 119307 19307 19307 19307 19307 19307 19307 1572 2502 2702 2702 2702 2702 2702 37950 48974 72928 8729 8729 8729 8729 8729 8729 872
9%; Score 18; DB 4; ).0%; Pred. No. 4.6; ) 0; Mismatches 86	US/09134001C US/09134001C CACID AND AMINO ACI CACID AND AMINO ACID CACID AND A	-09-461-697-359 Sequence -09-38-907-183 Sequence -09-38-907-183 Sequence -09-218-207-183 Sequence -09-218-207-183 Sequence -09-09-913-1 Sequence -09-453-702B-57 Sequence -09-453-702B-57 Sequence -08-368-803-21 Sequence -08-578-096A-22 Sequence -09-08-578-096A-22 Sequence -08-578-096A-22 Sequence -09-08-578-096A-22 Sequence -09-08-578-096A-22 Sequence -09-08-578-096A-22 Sequence	US-08-985-916- US-08-985-916- US-08-985-916- US-08-985-022A US-09-427-048A US-09-094-287- US-09-094-287- US-09-612-964- US-09-612-964- US-09-612-964- US-09-612-964- US-09-612-964- US-09-612-964- US-09-612-964- US-09-612-964- US-09-612-964- US-09-18-3702B US-09-18-3702B US-09-18-3702B US-09-18-3702B US-09-18-3702B US-09-18-3702B US-09-18-3702B US-09-18-3702B US-08-578-0963
; Length 1587; 0; Indels 0;	THER		
Gaps 0;	G TO STAPHYLOCOCC		4, Appli 6, Appli 10, Appli 10, Appli 11, Appli 11, Appli 11, Appli 12, Appli 12, Appli 13, Appli 13, Appli 13, Appli 13, Appli 16, Appli 17, Appli 17, Appli 17, Appli 17, Appli 17, Appli 17, Appli 17, Appli 17, Appli 18, Appli 17, Appli 17, Appli 18, Appli 17, Appli 18, Appli 17, Appli 18, Appli 19, Appli 11, Appli 11, Appli 11, Appli 12, Appli 12, Appli 13, Appli 14, Appli 15, Appli 16, Appli 17, Appli 18, Appli 19, Appli 11, Appli 11, Appli 11, Appli 12, Appli 12, Appli 13, Appli 14, Appli 15, Appli 16, Appli 17, Appli 18, Appli 18, Appli 19, Appli 19, Appli 19, Appli 19, Appli 19, Appli 19, Appli 10, Appli 11, Appli 11, Appli 11, Appli 12, Appli 12, Appli 13, Appli 14, Appli 15, Appli 16, Appli 17, Appli 18, Appli 18, Appli 19, Appli 19, Appli 19, Appli 10, Appli 10, Appli 11, Appli 11, Appli 12, Appli 12, Appli 13, Appli 14, Appli 15, Appli 16, Appli 17, Appli 18, Appl

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TYPE: DNA
; ORGANISM: Staphylococcus epidermidis
US-09-134-001C-263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; ORGANISM: Staphylococcus epidermidis US-09-134-001C-2083
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US-09-134-001C-263/c
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: EDIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC-007
CURRENT APPLICATION NUMBER: US/09/134,001C
CURRENT FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR TILING DATE: 1997-10-08
PRIOR FILING DATE: 1997-08-14
SEQ ID NO 263
LENGTH: 393
LENGTH: 393
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 627, Application US/09221017B Patent No. 6444799 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 6380370
GENERAL INFORMATION:
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                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: PP1182
FILING DATE: 31-DEC-1997
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                       ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: ROSS, BIUGE C.
TITLE OF INVENTION: P. GINGIVALIS NUCLEOTIDES AND USES THEREOF NUMBER OF SEQUENCES: 1120
CORRESPONDENCE ADDRESS:
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APPLICATION NUMBER: PP1546 FILING DATE: 30-JAN-1998
                                                                                                            APPLICATION NUMBER: US/0 FILING DATE: 23-DEC-1998 CLASSIFICATION:
                                                                                                                                                                                                    COMPUTER: IBM Compatible OPERATING SYSTEM: Windows
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       y Match 4.9%; So Local Similarity 100.0%; P Conservative 0;
                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                      STREET:
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                                                                                                                                                                                                                                                                                                                                  Palo Alto
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Pred. No.
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Pred. No.
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US-08-532-828B-11/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 11, Application US/08532828B Patent No. 5688671
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 17; Conserv
                         APPLICATION NUMBER: US/08/532,828B
FILING DATE: 27-OCT-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 5-101450
FILING DATE: 27-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: PC-DOS/MS-I
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: SUZUKI, Tomoko
APPLICANT: TANAKA, Akiko
APPLICANT: MATSUI, Hiroshi
TITLE OF INVENTION: MUTANT ASPARTOKINASE GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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HYPOTHETICAL:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: SUGIMOTO, Masakazu
APPLICANT: OGAWA, Yuri
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      104 AAGTCCTGTGCATCATC 120
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REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: doub
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: PCT/
FILING DATE: 10-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    E: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C. 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
                                                                                                                                                                                                                                                                                                                                                                                    IBM PC compatible SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UNKNOWN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-APR-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.6%; Score 17;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PP2911
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24,618
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15;
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Sequence 12, Application US/08532828B

Patent No. 5688671

GENERAL INFORMATION:
APPLICANT: SUGIMOTO, Masakazu
APPLICANT: OGAWA, Yuri
APPLICANT: SUZUKI, Tomoko
APPLICANT: TANAKA, Akiko
APPLICANT: MATSUI, Hiroshi
TITLE OF INVENTION: MUTANT ASPARTOKINASE GENE
NUMBER OF SEQUENCES: 24
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Best Local Similarity
Matches 17; Conserv
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO: 11:
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STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: genomic DI
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                TELEFAX: 703-413-2220 INFORMATION FOR SEQ ID NO: 12:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 1263 nucleotides
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                                                                                                                                                                                                                                         APPLICATION NUMBER: JP 5:
FILING DATE: 27-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: MS-DOS TEXT EDITOR CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 1263 nucleotides
TYPE: nucleic acid
                                                                                                                                                                   REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMAÇION:
TELEPHONE: 703-413-3000
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   HYPOTHETICAL:
                   MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/532,828B FILING DATE: 27-OCT-1995 CLASSIFICATION: 435
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ZIP: 22202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: ARLINGTON
                                                                                                                                    TELEPHONE: //3-2220
TO NO:
                                                                                                                                                                                                                        NAME: NORMAN F. OBLON REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
                                      TOPOLOGY:
                                                      STRANDEDNESS: double
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                                      linear
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NO
: genomic DNA
NO
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ORIGINAL SOURCE:
ORGANISM: Corynebacterium glutamicum
STRAIN: AJ3463
US-08-532-828B-12
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US-08-532-828B-1/c
                                                           US-08-532-828B-1
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Matches
           Query Match
Best Local Similarity
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                                                                                                                                                                                                                        TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1643 nucleotides
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C.
STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-I
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: MATSUI, HIROSHI
TITLE OF INVENTION: MUTANT ASPARTOKINASE GENE
                                                                          ANTI-SENSE: NO
ORIGINIAL SOURCE:
ORGANISM: Corynebacterium glutamicum
STRAIN: ATCC 13869
                                                                                                                                          MOLECULE TYPE: ge
                                                                                                                                                                                                                                                                                         NAME: NORMAN F. OBLON
REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703,413-3000
                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 27-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 5-101450
FILING DATE: 27-APR-1993
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                                                                                                                                        HYPOTHETICAL:
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hes 17; Conservative
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CITY: ARLINGTON
STATE: VA
                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/532,828B FILING DATE: 27-OCT-1995
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                                                                                                                                                                                                                                                            TELEPHONE: / / TELEPHONE: / 703-413-2220
                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                         TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
17;
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OGAWA, Yuri
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUZUKI, Tomoko
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                            linear
                                                                                                                                                            genomic DNA
                                                                                                                                                                                             double
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           4.6%; Score 17; DB 1; 100.0%; Pred. No. 15;
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0;
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Mismatches
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0
                                Length 1643;
Indels
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161 CAAAGTGCCGGGATCAT 177

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936 CAAAGTGCCGGGATCAT 920

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US-08-532-828B-7/c
                                                                                                                           RESULT 9
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US-08-532-828B-2
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                                                                         Sequence 7, Application US/08532828B Patent No. 5688671
                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                Query Match
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                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: GER
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
APPLICANT:
                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 5-101450
FILING DATE: 27-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: PC-DOS/MS-I
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: SUGIMOTO, Masakazu
APPLICANT: OGAWA, Yuri
APPLICANT: SUZUKI, Tomoko
APPLICANT: TANAKA, Akiko
APPLICANT: MATSUI, Hiroshi
TITLE OF INVENITON: MUTANT ASPARTOKINASE GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
                                                                                                                                                                                         161 CANAGTGCCGGGATCAT 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                     936 CAAAGTGCCGGGATCAT 920
                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: lin
                                                                                                                                                                                                                                                                                                                           ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/532,828B FILING DATE: 27-OCT-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: ARLINGTON
STATE: VA
                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
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                                                                                                                                                                                                                                  17; Conserv

    Application US/08532828B
    5688671

                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 1643 nucleotides
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                                                                                                                                                                                                                                 Conservative
TANAKA, Akiko
                           SUGIMOTO, Masakazu
OGAWA, Yuri
           SUZUKI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                         Corynebacterium glutamicum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                       linear
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                                                                                                                                                                                                                                                                                                                                                                                        genomic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                       double
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ative 0;
                Tomoko
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                                                                                                                                                                                                                                                                 4.68;
                                                                                                                                                                                                                                               Score 17;
Pred. No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-764-0 PCT
                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                 No.
                                                                                                                                                                                                                                              DB 1; Length 1643; 15;
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                                                                                                                                                                                                                               Gaps
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RESULT 10
US-08-532-828B-8/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-532-828B-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 703-413-2220 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                            APPLICANT: TANAKA, ANALA APPLICANT: MATSUI, HIROShi
APPLICANT: MATSUI, HIROShi
TITLE OF INVENTION: MUTANT ASPARTOKINASE GENE
TITLE OF STOTTENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: JP 5-101450 FILING DATE: 27-APR-1993 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 1643 nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 27-OCT-CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: PC-DOS/MS-I
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HYPOTHETICAL: NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 936 CARAGTGCCGGGATCAT 920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        161 CAAAGTGCCGGGATCAT 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: MATSUI, HIROSHI TITLE OF INVENTION: MUTANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY:
LOCATION:
COUNTRY:
                                                                                       CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 100.0%; In the second secon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Corynebacterium glutamicum STRAIN: ATCC13869
                                                   STATE:
                                                                                                                                          STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: /UJ TELEPH
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                                                                                                                                                                                              ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER:
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                                                                                                                                E: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C. 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
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USA
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IBM PC compatible
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JEFFERSON DAVIS HIGHWAY, SUITE 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MUTANT ASPARTOKINASE GENE
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100.0%; Pred. No.
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ZIP: 22202 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy

IBM PC compatible SYSTEM: PC-DOS/MS-DOS

Floppy disk

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US-08-532-828B-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 11
US-08-532-828B-9/c
                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: SUGIMOTO, MAS
APPLICANT: OGAWA, YUri
APPLICANT: SUZUKI, TOMOK
APPLICANT: TANAKA, AKIKC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 4.6%; Score 17; Best Local Similarity 100.0%; Pred. No. Matches 17; Conservative 0; Mismatcl
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9, Application US/08532828B Patent No. 5688671
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APPLICATION NUMBER: JP 5-101450
FILING DATE: 27-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10-764-0
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/532,828B
FILING DATE: 27-OCT-1995
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TELEPHONE: 703-413-3000
TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORIGINAL SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: 96
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                            APPLICANT: MATSUI, HIROSHI
TITLE OF INVENTION: MUTANT ASPARTOKINASE GENE
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      161 CAAAGTGCCGGGATCAT 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 936 CAMAGTGCCGGGATCAT 920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/532,828B FILING DATE: 27-OCT-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: PC-DOS/MS-I
SOFTWARE: MS-DOS TEXT EDITOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Corynebacterium glutamicum STRAIN: AJ3463
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                                                                                                                                                                                                             STREET: 1755 S.
CITY: ARLINGTON
                                                                                                                                                               COUNTRY: U
ZIP: 22202
                                                                                                                                                                                                                                          ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C. STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
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                                                                                                                                                                                                                                                                                                                                                                                                                      SUGIMOTO, Masakazu
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US-08-532-828B-9
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US-08-532-828B-10/c
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Best Local Similarity
Matches 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION NUMBER: JP 5-1
APPLICATION NUMBER: JP 5-1
FILING DATE: 27-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
                                                APPLICATION NUMBER: US/08/532,828B
FILING DATE: 27-OCT-1995
CLASSIFICATION : 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 5-101450
FILING DATE: 27-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FOR PC PODS/MS-DOS
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: SUGIMOTO, MASAKAZU
APPLICANT: OGAWA, YUTİ
APPLICANT: SUZUKI, TOMOKO
APPLICANT: TANAKA, AKİKO
APPLICANT: MATSUI, HİTOShİ
TITLE OF INVENTION: MUTANT ASPARTOKINASE GENE
NUMBER OF SEQUENCES: 24
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TELECOMMUNICATION INFORMATION:
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ZIP: 22202
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                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C. STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
                 REGISTRATION NUMBER: 24,618 REFERENCE/DOCKET NUMBER: 10
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                   10-764-0 PCT
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TELEPHONE:

703-413-3000

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us-08-700-359-7/c
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                                                                            TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP-350
FILING DATE: 04-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1643 nucleotid
 MOLECULE TYPE:
                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: PC-DOS/MS-I
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: SUZUKI, TOMOKO
APPLICANT: TANAKA, AKIKO
APPLICANT: MATSUI, HIROSHI
TITLE OF INVENTION: METHOD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: SUGIMOTO, MASAKAZU APPLICANT: USUDA, YOSHIHIRO APPLICANT: SUZUKI, TOMOKO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE
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HYPOTHETICAL:
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                  TOPOLOGY:
                            STRANDEDNESS:
                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/700,359 FILING DATE: 08-OCT-1996 CLASSIFICATION: 435
                                                                                                                                                           REFERENCE/DOCKET NUMBER:
                                                                                                                                                                            REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                       COMPUTER:
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                                                                LENGTH:
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                                             nucleic acid
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SYSTEM: PC-DOS/MS-DOS
                linear
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                                                                                                                                                                                                                                           JP-35019
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100.0%; Pred. No.
                                                                                                                                                                         24,618
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                                                                                                                                                           10-819-0 PCT
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US-08-700-359-8
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US-08-700-359-8/c
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US-08-700-359-7
               Best Local
      Matches
                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 8, Application US/08700359 Patent No. 5766925
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                            MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 04-MAR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: MS-DOS TEXT EDITOR CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORIGINAL SOURCE:
ORGANISM: Corynebacterium glutamicum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ANDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: TANAKA, AKIKO
APPLICANT: MATSUI, HIROSHI
TITLE OF INVENTION: METHOD OF PRODUCING L-LYSINE
NUMBER OF SEQUENCES: 22
                                                                                        NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        161 CAAAGTGCCGGGATCAT 177
                                                                         IDENTIFICATION METHOD:
                                                                                                                                 ORGANISM: Corynebacterium glutamicum STRAIN: ATCC13869
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                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 24
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                    NAME: NORMAN F. OBLON
                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: 08-OC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: ARLINGTON
STATE: VA
                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
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                  Similarity
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                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                          genomic DNA
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100.0%; Pr
0;
               4.6%; Score 17; DB 1; 100.0%; Pred. No. 15;
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                                                                                                                                                                                                                                                                                                                                 24,618
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Pred. No.
   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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                           Length 1643;
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0; Gaps
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161 CAAAGTGCCGGGATCAT 177

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936 CAAAGTGCCGGGATCAT 920

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US-08-700-359-10/c
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                                                                   RESULT 16
US-08-674-168-21/c
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: SUZUKI, ...
APPLICANT: TANAKA, AKIKO
APPLICANT: TANAKA, AKIKO
APPLICANT: MATSUI, HIROSHI
TITLE OF INVENTION: METHOD OF PRODUCING L-LYSINE
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIYAK, MCCLELLAND, MAIER & NEUSTADT, P.C.
ADDRESSEE: OBLON, SPIYAK, MCCLELLAND, MAIER & NEUSTADT, P.C.
               Sequence 21, Application US/08674168 Patent No. 5804414 GENERAL INFORMATION:
                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:

APPLICATION NUMBER: JP-35019

FILING DATE: 04-MAR-1994

ATTORNEY/AGENT INFORMATION:

NAME: NORMAN F. OBLON

REGISTRATION NUMBER: 24,618

REFERENCE/DOCKET NUMBER: 10-81

TELECOMMUNICATION INFORMATION:

TELEPHONE: 703-413-3300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 703-413-2220 INFORMATION FOR SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS-DOS TEXT EDITOR
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: SUGIMOTO, MASAKAZU APPLICANT: USUDA, YOSHIHIRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
APPLICANT: MORIYA, Mika
                                                                                                                                                936 CAAAGTGCCGGGATCAT 920
                                                                                                                                                                                     161 CAAAGTGCCGGGATCAT 177
                                                                                                                                                                                                                                                                                                                                   NAME/KEY:
                                                                                                                                                                                                                          Local Similarity hes 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/700,359 FILING DATE: 08-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: doub
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                          STRAIN:
                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM:
                                                                                                                                                                                                                                                                                                                     LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22202
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ATCC13869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1643
                                                                                                                                                                                                                       4.6%; Score 17; DB 1; llarity 100.0%; Pred. No. 15; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                   mat peptide
964..1479
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  genomic DNA
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                                                                                                                                                                                                                            0
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                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                            0;
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; ORGANISM: Brevibac; STRAIN: ATCC 13869 US-08-674-168-21
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US-08-596-366-5/c
                                                                                                                                                                          Sequence 5, Application US/08596366 Patent No. 5876983
                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 413-3000
TELEFAX: (703) 413-2220
TELEX: 24885 OPAT UR
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: METHOD OF AMPLIFYING GENE USING TITLE OF INVENTION: ARTIFICIAL TRANSPOSON NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: OPION 7--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: JP 7-166541 FILING DATE: 30-JUN-1995 ATTORNEY/AGENT INFORMATION: NAME: OBLON, NORMAN F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
           APPLICANT: SUGIMOTO, MASAKAZU
APPLICANT: SUZUKI, TOMOKO
APPLICANT: MATSUI, HICOSHI
APPLICANT: IZUI, Katsura
TITLE OF INVENTION: MUTANT PHOSPHOENOLPYRUVATE CARBOXYLASE,
TITLE OF INVENTION: ITS GENE, AND PRODUCTION METHOD OF AMINO ACID
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                          161 CAAAGTGCCGGGATCAT 177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: ARLINGTON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: OBLON, NORMAN F.
REGISTRATION NUMBER: 24,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 01 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
                                                                                                                                                                                                                                                                                                                                                          y Match 4.6%; So
Local Similarity 100.0%; F
hes 17; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/674,168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1643 base pairs
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YOKOZEKI, Kenzo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HAYAKAWA, Atsushi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patentin Release #1.0, Version #1.30
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OBLON, SPIVAK, MCCLELLAND, MAIER, & NEUSTADT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JUL-1996
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                                                                                                                                                                                                                                                                                                                                                                                Score 17;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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RESULT 18
US-08-596-366-7/c
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                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: SUGIMOT,
APPLICANT: SUZUKI,
APPLICANT: MATSUI,
APPLICANT: IZUI, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 4.6%; Score 17; DB 2; Length 1643; Best Local Similarity 100.0%; Pred. No. 15; Matches 17; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                           Sequence 7, Application US/08596366 Patent No. 5876983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (703) 413-3000
TELEFAX: (703) 413-2220
TELEX: 248855 OPAT UR
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: JE
FILING DATE: 24-AUG-19
PRIOR APPLICATION DATA:
                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                     TITLE OF INVENTION: MUTANT PHOSPHOENOLPYRUVATE CARBOXYLASE,
TITLE OF INVENTION: ITS GENE, AND PRODUCTION METHOD OF AMINO ACID
NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                    161 CAAAGTGCCGGGATCAT 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: OBLON, NORMAN F.
REGISTRATION NUMBER: 24,618
REFERENCE, DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                   936 CAAAGTGCCGGGATCAT 920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: JP 6-153876
FILING DATE: 05-JUL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 24-AUG-
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 1/J
STREET: ARLINGTON
CITY: ARLINGTON
CTATE: VIRGINIA
ADDRESSEE: OBLUN, SELL.....
ADDRESSEE: P.C.
ADDRESSEE: 1755 JEFFERSON DAVIS HIGHWAY, SUITE # 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN: ATCC13869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
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FILING DATE: 29-APR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic acid
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                                                                                                                                                                                  SUZUKI, Tomoko
MATSUI, Hiroshi
                                                                                                                                                             IZUI, Katsura
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                                                                                                                                                                                                                        SUGIMOTO, Masakazu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mat_peptide
217..1482
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
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SYSTEM: PC-DOS/MS-DOS
                                                                 OBLON, SPIVAK, McCLELLAND, MAIER, & NEUSTADT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA (genomic)
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N: 435
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US-08-967-104-5/c
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                                                                                                                                                                                                                                                                                           Sequence 5, Application US/08967104 Patent No. 5919694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: SUGIMO
APPLICANT: SUZUKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (703) 413-3000
TELEFAX: (703) 413-2220
TELEX: 24885 OPAT UR
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                  TITLE OF INVENTION: MUTANT PHOSPHOENOLPYRUVATE CARBOXYLASE,
TITLE OF INVENTION: ITS GENE, AND PRODUCTION METHOD OF AMINO ACID
NUMBER OF SEQUENCES: 12
                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN: ATCC13869 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: JP 6-153876
FILING DATE: 05-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: OBLON, NORMAN F.
REGISTRATION NUMBER: 24,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: DNA (genomic) ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 24,618
REFERENCE/DOCKET NUMBER: 10-784-0 PCT
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 24-AUG-1993
PRIOR APPLICATION DATA;
APPLICATION NUMBER: JP 5-
FILING DATE: 24-AUG-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                  936 CAAAGTGCCGGGATCAT 920
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APPLICATION NUMBER:
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MEDIUM TYPE: Floppy disk
                                   STREET: 1755 JEI
CITY: ARLINGTON
STATE: VIRGINIA
COUNTRY: UZIP: 22202
                                                                                       ADDRESSEE:
                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity es 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/596,366 FILING DATE: 29-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 22202
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: VIRGINIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleic acid
                                                                1755 JEFFERSON DAVIS HIGHWAY, SUITE # 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1643 base pairs
                                                                                                                                                                                                                SUGIMOTO, Masakazu
SUZUKI, Tomoko
MATSUI, Hiroshi
                                                                                                                                                                                             IZUI, Katsura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.6%; Score 17; DB 2; ilarity 100.0%; Pred. No. 15; Conservative 0; Mismatches
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964..1482
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                                                                                                    OBLON, SPIVAK, McCLELLAND, MAIER, & NEUSTADT
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SYSTEM: PC-DOS/MS-DOS
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COMPUTER READABLE FORM:

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FILING DATE: 24-AUG-193

PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 6-1538;
APPLICATION NUMBER: JP 6-1538;
APPLICATION NUMBER: JP 6-1538;
APPLICATION NUMBER: JP 6-1538;
APPLICATION NORMAN F.
REGISTRATION NUMBER: 24,618
REGISTRATION NUMBER: 10-7;
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 413-3000
TELEX: 24885 OPAT UR
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                              Sequence 7, Application US/08967104
Patent No. 5919694
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 4.6%;
Best Local Similarity 100.0%;
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PRIOR APPLICATION DATA:
                                                                                                                                             APPLICANT: SUGIMOTO, MASAKAZU
APPLICANT: SUZUKI, TOMOKO
APPLICANT: MATSUI, HICOSHI
APPLICANT: MATSUI, HICOSHI
APPLICANT: IZUI, Katsura
TITLE OF INVENTION: MUTANT PHOSPHOENOLPYRUVATE CARBOXYLASE,
TITLE OF INVENTION: ITS GENE, AND PRODUCTION METHOD OF AMINO ACID
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
ORIGINAL SOURCE:
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COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/596,366 FILING DATE: 29-APR-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Corynebacterium glutamicum STRAIN: ATCC13869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                     CITY: ARLINGTON
STATE: VIRGINIA
                                                                                       STREET:
                             COUNTRY:
                                                                                                             ADDRESSEE:
                                                                                                                               ADDRESSEE:
                   22202
                                                                                         E: P.C.
1755 JEFFERSON DAVIS HIGHWAY, SUITE # 400
                                   USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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SYSTEM: PC-DOS/MS-DOS
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217..1482
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                                                                                                                             OBLON, SPIVAK, McCLELLAND, MAIER, & NEUSTADT
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24-AUG-1993
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                             Sequence 3, Application US/08985908
Patent No. 6004773
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 6-1
FILING DATE: 05-UUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: OBLON, NORMAN F.
REGISTRATION UNUMBER: 24.61
REFERENCE/DOCKET NUMBER: 1
COMPUTER READABLE FORM:
MEDIUM TYPE: FLORPY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/985,908
FILING DATE: 05-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (703) 413-3000
TELEFAX: (703) 413-2220
TELEX: 24885 OPAT UR
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                 APPLICANT: MASAYUKI ARAKI, MASAKAZU SUGIMOTO, YASUHIKO YOSHIHARA, AND TSUYOSHI
TITLE OF INVENTION: METHOD FOR PRODUCING L-LYSINE
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
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ORIGINAL SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                               936 CAAAGTGCCGGGATCAT 920
                                                                                                                                        Clir.
COUNTRY: UC.
22202
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 1643 base pairs
                                                                                                                                                                                                  STREET:
                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid
                                                                                                                                                                                      ARLINGTON
                                                                                                                                                                                                      E: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C. 1755 S. JEFFERSON DAVIS HIGHWAY, FOURTH FLOOR
                                                                                                                                                                     USA
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964..1482
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-APR-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.6%; Score 17; DB 2; 100.0%; Pred. No. 15; tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JP 5-209776
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JP 6-153876
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                                                        Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24,618
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CLASSIFICATION:

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATE: 05-DEC-LIVE ATTORNEY/AGENT INFORMATION: NAME: NORMAN F. OBLON REGISTRATION NUMBER: 24,618
TELECOMMUNICATION INFORMATION: 703-413-3000
                                                                                                                                                TELEFAX: 703-413-2220 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 1643 bases
TYPE: nucleic acid
                             MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                              NAME: NORMAN F. OBLON
REGISTRATION NUMBER: 24,618
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 8-325659
FILING DATE: 05-DEC-1996
                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                               FILING DATE: 05-DEC-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: genomic DNA ORIGINAL SOURCE:
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                                                                                                                                               TELEPHONE: 703-413-2220
                 ORGANISM:
                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/985,908 FILING DATE: 05-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C. STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, FOURTH FLOOR CITY: ARLINGTON
                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE:
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ATCC 13869
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MASAYUKI ARAKI, MASAKAZU SUGIMOTO, YASUHIKO YOSHIHARA, AND TSUYOSHI NA
IVENTION: METHOD FOR PRODUCING L-LYSINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4.6%; Score 17; ilarity 100.0%; pred. No. Conservative 0; Mismatc
          Brevibacterium lactofermentum
                                                              linear
                                                                                                                                                                                                                                                                                                                                                                                              IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                 PatentIn
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                                             genomic DNA
                                                                                double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   double
                                                                                                                                                                                                                                                                                                                                                                              Release #1.0, Version #1.30
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                                           Query Match
Best Local Similarity 100.
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INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
1.F.NGTH: 1643 bases
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                         FILING DATE: 05-DEC-1996
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
REGISTRATION UNMBER: 24,618
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
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                                                                                                                                                                                                 MOLECULE TYPE: genomic DNA ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION NUMBER:
                                                                                                                                                       FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Rela
936 CAAAGTGCCGGGATCAT 920
                            161 CAAAGTGCCGGGATCAT 177
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MEDIUM TYPE: Floppy disk
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TITLE OF INVENTION: METHOD FOR PRODUCING L-LYSINE
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS
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                                                                                                                                          NAME/KEY:
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                                                                                                                              LOCATION:
                                                                                                                                                                       STRAIN: ATCC 13869
                                                                                                                                                                                      ORGANISM:
                                                                                                                                                                                                                                                      TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: 05-DE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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les 17; Conserv
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217..1482
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05-DEC-1996
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100.0%; Pred. No.
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                                                                                 DB 3;
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15;
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US-08-852-730-12/c ; Sequence 12, Application US/08852730

RESULT 24

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RESULT 25
US-08-852-730-13/c
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Best Local Similarity
Matches 17; Conserva
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                                                                                                                                                                                                                                                                                                                                                     Sequence 13, Application US/08852730 Patent No. 6090597
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO: 12:
                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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APPLICANT: MASAKO IZUI, ATSUSHI HAYAKAWA, YASUHIKO YOSHIHARA, AND TSUYOSHI
APPLICANT: NAKAWATSU
TITLE OF INVENTION: METHOD OF PRODUCING L-LYSINE
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: JP 8-142812
FILING DATE: 05-JUN-1996
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
REGISTRATION NUMBER: 24,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 05-07-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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LENGTH: 1643 base pairs
TYPE: nucleic acid
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ORIGINAL SOURCE:
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STATE: VA
STATE: 22026
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSE: OBLON, SPIVAK, MCCLELLAND, MAIER AND NEUSTADT, P.C. STREET: 1755 S. JEFFERSON DAVIS HIGHWAY
CITY: ARLINGTON
                                                                                                                                                                                                                                                                     APPLICANT: SEIKO HIRANO, MASAKAZU SUGIMOTO, EIICHI NAKANO, APPLICANT: MASAKO IZUI, ATSUSHI HAYAKAWA, YASUHIKO YOSHIHARA, AND TSUYOSHI APPLICANT: NAKAMATSU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      161 CAAAGTGCCGGGATCAT 177
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STATE: Va
22026
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Brevibacterium lactofermentum STRAIN: ATCC 13869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Genomic DNA
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                                                                                                                                                                                                                                                    METHOD OF PRODUCING L-LYSINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
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GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 8-142812
FILING DATE: 05-JUN-1996
ATTORNEY/AGENT INFORMATION:
NAME: NORMAN F. OBLON
REGISTRATION NUMBER: 24 618
TELECOMMUNICATION INFORMATION:
                   • TELEFAX: 703-413-2220 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 1643 base pair
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ATTORNEY/AGENT INFORMATION:
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ORIGINAL SOURCE:
ORGANISM: Brev
                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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SEQUENCE CHARACTERISTICS:
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STATE: VA
TTD: 22026
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FILING DATE: 05-07-1997
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LOCATION:
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                   TELEPHONE: /vv
TELEPHONE: /vv
TO3-413-2220
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                                                                                                                                                                                                    FILING DATE: 0:
CLASSIFICATION:
                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/852,730 FILING DATE: 05-07-1997
                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER AND NEUSTADT, P.C. STREET: 1755 S. JEFFERSON DAVIS HIGHWAY
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217..1482
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                                                      703-413-3000
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                                                         Matches
                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 3, Application US/08985916 Patent No. 6221636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                    TELEFAX: 703-413-2220
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: ATSUSH
                                                                                                                           MOLECULE TYPE: genomoriginal Source:
ORGANISM: Brevibact
STRAIN: ATCC 13869
                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 24,618
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-413-3000
                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: JP 8-325658
FILING DATE: 05-DEC-1996
ATTORNEY_AGENT INFORWATION:
NAME: NORMAN F. OBLON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
936 CAAAGTGCCGGGATCAT 920
                           161 CAAAGTGCCGGGATCAT 177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
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ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           161 CAAAGTGCCGGGATCAT 177
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                                                                     Local
                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 05-DE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                              LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
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LOCATION:
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                                                                   Similarity
                                                                                                                                                                                                              nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1: 1755 S. JEFFERSON DAVIS HIGHWAY, FOURTH FLOOR ARLINGTON
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                                                                                                                                                                                                                                1643 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                M: Brevibacterium lactofermentum
ATCC 13869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1643 base pairs
                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATSUSHI HAYAKAWA, MASAKAZU SUGIMOTO, YASUHIKO YOSHIHARA, AND TSUYOSHI
VENTION: METHOD FOR PRODUCING L-LYSINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                        PatentIn Release #1.0, Version #1.30
                                                                                                                                     Brevibacterium lactofermentum
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964..1482
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                                                                                                                                                                       genomic DNA
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                                                                     100.0%;
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L00.0%; Pred. No.
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                                                    0;
                                                                   Score 17; 
; Pred. No.
                                                    Mismatches
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                                                                  DB 4;
15;
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                                                                              Length 1643;
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US-08-985-916-4/c
                                                                                                                                                                             Sequence 6, Application US/08985916 Patent No. 6221636 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Patent No. 6221636
GENERAL INFORMATION:
APPLICANT: ATSUSHI HAYAKAWA, MAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 703-413-2220 INFORMATION FOR SEQ ID NO:
                                                                                                    APPLICANT: ATSUSHI HAYAKAWA, MASAKAZU SUGIMOTO, YASUHIKO YOSHIHARA, AND TSUYOSH TITLE OF INVENTION: METHOD FOR PRODUCING L-LYSINE NUMBER OF SEQUENCE: 24
CORRESPONDENCE ADDRESS:
COMPUTER READABLE FORM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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ORIGINAL SOURCE:
ORGANISM: Bre
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CURRENT APPLICATION DATA:
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LENGTH: 1643 bases
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FILING DATE: 05-DEC-1996
ATTORNEY/AGENT INFORMATION:
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CORRESPONDENCE ADDRESS:
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                                              STREET: 1755 S. CITY: ARLINGTON
                                COUNTRY:
                                                         ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTAD' STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, FOURTH FLOOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: 703-413-2220
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN: ATCC 13869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: 05-DECCLASSIFICATION: 439
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY:
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                   22152
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                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       genomic DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JEFFERSON DAVIS HIGHWAY, FOURTH FLOOR
                                                                                                                                                                                                                                                                                                                                                                                                      4.6%; Score 17; DB 4; 100.0%; Pred. No. 15;
                                                                                 SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US/08/985,916
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                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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Floppy disk

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; NAME/KEY:
; LOCATION:
US-08-985-916-6
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Best Local Similarity
Matches 17; Conserv
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GENERAL INFORMATION:
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APPLICATION NUMBER: JP 8-325658

FILING DATE: 05-DEC-1996

ATTORNEY/AGENT INFORMATION:

NAME: NORMAN F. OBLON

REGISTRATION NUMBER: 24,618

TELECOMMUNICATION INFORMATION:

TELEPHONE: 703-413-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 1643 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: genomic DNA ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Trustees of the University of Pennsylva APPLICANT: Wilson, James M. APPLICANT: Fisher, Krishna J. APPLICANT: Chen, Shu-Jen APPLICANT: Weitzman, Matthew TITLE OF INVENTION: Improved Adenovirus Virus and NUMBER OF SEQUENCES: 10
                    CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 08/331,381
FILING DATE: 28-OCT-1994
                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               936 CAAAGTGCCGGGATCAT 920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      161 CAAAGTGCCGGGATCAT 177
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
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COMPUTER: IF
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STRANDEDNESS: double
TOPOLOGY: linear
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                                                                                               APPLICATION NUMBER: FILING DATE:
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                                                                                                                                                                                                                MEDIUM TYPE:
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                                                                                                                                                                                                                                                       19477
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964..1482
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                              Floppy disk
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                                                                                                                     US/08/836,022A
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US-08-836-022A-10
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US-09-427-048A-10/c
US-09-427-048A-10
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GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 10
SEQUENCE CHARACTERISTICS:
LENGTH: 19307 base pairs
TYPE: nucleic acid
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9200
TELEPHAX: 215-540-5818
                                                                                                                              INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Bak, Mary E. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOETWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Weitzman, Matthew
TITLE OF INVENTION: Improved Adenovirus Virus and
Methods of Use Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Trustees of the University of Pennsylvania Wilson, James M. Fisher, Krishna J.
                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,215
                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/836,022
FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Howson and Howson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                  REFERENCE/DOCKET NUMBER: GNVPN.008PCT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9200
TELEFAX: 215-540-5818
                                   TOPOLOGY: unknown MOLECULE TYPE: cDNA
                 SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/427,048A FILING DATE: 21-Oct-1999 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Spring House
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
                                                                       LENGTH: 19307 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/09427048A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 19477
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100.08; Pr
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100.0%; Pred. No.
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                                                                                    Sequence 1, Application US/09094287
PATENT NO. 6084071
GENERAL INFORMATION:
APPLICANT: Racie, Lisa
APPLICANT: Jacobs, Kenneth
                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 617-876-5851 INFORMATION FOR SEQ ID NO: 8:
                 APPLICANT: Jacobs, Kenneth
TITLE OF INVENTION: Human L105 Proteins and Polynucleotide
TITLE OF INVENTION: Encoding Same
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 535 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/094,287
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TITLE OF INVENTION: Human L105 Proteins and Polynucleotide
TITLE OF INVENTION: Encoding Same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                          256 TGCATCAGCTGCTGCA 241
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                                                                                                                                                                                                                                                                                                                                                                                           DESCRIPTION: /desc = "EST sequence"
     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
REFERENCE/DOCKET NUMBER: GI
                                                                                                                                                                                                                                                                                                              Local Similarity
les 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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87 CambridgePark Drive
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617-876-5851
                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
Genetics Institute, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                double
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100.08; Pr
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100.0%; Pred. No.
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100.0%; Pred. No. 15;
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US-09-094-287-3/c
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                                                                                                                                                                                                                                                                                                        Sequence 3, Application US/09094287
Patent No. 6084071
GENERAL INFORMATION:
APPLICANT: Racie, Lisa
APPLICANT: Jacobs, Kenneth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY:
LOCATION:
FEATURE:
                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                             ADDRESSEE: Genetics Institute,
                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-498-8224
                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   342 TGCATCAGCTGCTGCA 327
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              APPLICATION NUMBER: US/09/094,287 FILING DATE: CLASSIFICATION:
                                                                                                                                                                                             STREET: 87 Cambr
CITY: Cambridge
                                                                                                                                                                      COUNTRY:
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EDNESS: double
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80..481
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149..481
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                                                                                                                                                                                                                                                                                       Human L105 Proteins and Polynucleotide
                                                                                                                                                                                                                                                                            Encoding Same
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100.0%; Pred. No.
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                                                                            Version #1.30
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NAME: Brown, Scott A REGISTRATION NUMBER:

Scott A.

32,724

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; LOCATION: (1)..(1569)
US-09-612-964-1
                                                                         RESULT 36
US-08-062-472B-2
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-498-8224
TELEPHONE: 617-876-5851
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 932 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/612,964
CURRENT FILING DATE: 2000-07-10
PRIOR APPLICATION NUMBER: RU 99114325
PRIOR FILING DATE: 199-07-09
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 1
LENGTH: 1572
                Sequence 2, Application US/08062472B Patent No. 5695954
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1, Application US/09612964 Patent No. 6403342
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Best Local Similarity
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                                                                                                                                                                                                                                     Matches
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APPLICANT: Lunts Maria Grigorievna
APPLICANT: Kozlov Yury Ivanovich
APPLICANT: Ivanovskaya Lirina Valerievna
APPLICANT: Voroshilova Elvira Borisovna
TITLE OF INVENTION: DNA CODING FOR MUTANT ISOPROPYLMALATE SYNTHASE,
TITLE OF INVENTION: L-LEUCINE-PRODUCING MICROORGANISM AND METHOD FOR PRODUCING
TITLE OF INVENTION: L-LEUCINE-PRODUCING MICROORGANISM AND METHOD FOR PRODUCING
FILE REFERENCE: 193845USO
                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Escherichia coli
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               269 TGCATCAGCTGCTGCA 284
                                                                                                                                                         547 ATCATCAGCGGCCTGT 562
                                                                                                                                                                              196 ATCATCAGCGGCCTGT 211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY:
                                                                                                                                                                                                                                   watch 4.3%; Score 16; DB Local Similarity 100.0%; Pred. No. 49 nes 16; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CDS
85..519
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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154..519
Sherwood, Nancy G M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
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; Pred. No. 49;
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                                                                                                                                                                                                                                                        DB 4; Length 1572; 49;
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APPLICANT: Hormas, Mary Beth
APPLICANT: Thomas, Mary Beth
APPLICANT: Thomas, Mary Beth
APPLICANT: Portbury, Stuart D.
APPLICANT: Portbury, Stuart D.
APPLICANT: Portbury, Stuart D.
APPLICANT: Rortnow, Compositions and Methods for Diagnosing
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING
TITLE OF INVENTION: CELL DEATH
FILLE REFERENCE: 10001-005-999
CURRENT APPLICATION NUMBER: US/09/461,697
CURRENT APPLICATION NUMBER: US/09/461,697
CURRENT FILING DATE: 1999-12-14
NUMBER OF SEQ ID NOS: 466
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 359
LENGTH: 2702
TYPE: DNA
ORGANISM: Homo sapiens
US-09-461-697-359
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US-09-461-697-359
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                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: COGENT NEUROSCIENCE, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 359, Application US/09461697 Patent No. 6277974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (503) 226-7391
TELEFAX: (503) 228-9446
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 2502 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Lescheid, David W
TITLE OF INVENTION: DNA ENCODING TWO FISH NEUROPEPTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: KLARQUIST, SPARKMAN, CAMPBELL, LEIGH &
ADDRESSEE: WHINSTON, LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 28107
TELECOMMUNICATION INFORMATION:
TELEPHONE: (503) 226-7391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 14-MAY-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1142 TTGCATCAGCTGCTGC 1157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: POLLEY, RICHARD J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                268 TIGCATCAGCIGCTGC 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA
ZIP: 97204-2988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: PORTLAND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: ONE WORLD TRADE CENTER, SUITE 1600, 121 S.W. STREET: SALMON STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OREGON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             McRory, John E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Parker, David B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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100.0%;
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                                                                               RESULT 39
US-09-338-907-183/c
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US-09-453-702B-258
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                              Sequence 183, Application US/09338907 Patent No. 6265546 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 258, Application US/09453702B Patent No. 6365723 GENERAL INFORMATION:
                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                     Query Match
APPLICANT: Cohen, Daniel APPLICANT: Blumenfeld, Marta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (608) 251-910 INFORMATION FOR SEQ ID NO: 258:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2500 AACTCAGTTTGCATCA 2515
                                                                                                                                                 371 CCCAAGAAAAGGGGAA 386
                                                                                                                                                                  311 CCCAAGAAAAGGGGAA 326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       260 AACTCAGTTTGCATCA 275
                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Seay, Nicholas J.
REGISTRATION NUMBER: 27386
REFERENCE/DOCKET NUMBER: 90
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 251-5000
                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
SEQUENCE DESCRIPTION: SEQ ID NO: 258:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/110,955
FILING DATE: 04-DEC-1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-005/MS-DOS SOFTWARE: Word Perfect 8.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: <Unknown> PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: No. 6365723el Sequences of E. coli 0157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Blattner, Frederick R.
                                                                                                                                                                                                                      16;
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                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/453,702B FILING DATE: 03-Dec-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 1 South Pinckney Street CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: US
ZIP: 53701-2113
                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 8729 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Quarles & Brady
                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Plunkett,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Burland,
                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.3%; Score 16;
100.0%; Pred. No.
                                                                                                                                                                                                                                                    4.38;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          251-9166
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                                                                                                                                                                                                                                    Score 16;
Pred. No.
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RESULT 40
US-09-218-207-183/c
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                                                                                          GENERAL INFORMATION:
APPLICANT: Cohen, Daniel
APPLICANT: Blumenfeld, Marta
APPLICANT: 11ya, Chumakov
APPLICANT: 11ya, Chumakov
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: Prostate cancer gene
FILE REFERENCE: GENSET.018CP1
                                                                                                                                                                                                                               Sequence 183, Application US/09218207 Patent No. 6346381
                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
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LENGTH: 37950
                  CURRENT APPLICATION NUMBER: US/09/218,207
CURRENT FILING DATE: 1998-12-22
EARLIER APPLICATION NUMBER: 08/996,306
EARLIER FILING DATE: 1997-12-22
  EARLIER FILING DATE: 1997-1: EARLIER APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EARLIER APPLICATION NUMBER: 08/996,306
EARLIER FILING DATE: 1997-12-22
EARLIER APPLICATION NUMBER: 60/099,658
EARLIER FILING DATE: 1998-09-09
EARLIER APPLICATION NUMBER: 09/218,207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/338,907 CURRENT FILING DATE: 1999-06-23
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APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: PROSTATE CANCER GENE
FILE REFERENCE: GENSET.18CP1CP
                                                                                                                                                                                                                                                                                                                                                34677 TCAGAAAGAAGTGATG 34662
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: exon
LOCATION: 25520..26016
OTHER INFORMATION: exon8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: exon7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: exon
LOCATION: 23387..23510
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LOCATION: 21789.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: exon6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: exon5
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LOCATION: 1982
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LOCATION: 1267
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                                                                                                                                                                                                                                                                                                                                                                                      142 TCAGAAAGAAGTGATG 157
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mes 16; Conserva
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100.0%; Pred. No.
60/099,658
                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 37950;
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                                                                                                                                                                                                                                                                                                                                                                                                                          0
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EARLIER FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 578
SOFTWARE: Patent.pm
SEQ ID NO 183
LENGTH: 37950

TYPE: DNA

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; SEQ ID NO 17
LENGTH: 48974
; TYPE: DNA
; ORGANISM: Mus musculus
US-08-920-422-17
                                 õ
Db 43070 TGTCAGAAAGAAGTGA 43085
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Db 34677 TCAGAAAGAAGTGATG 34662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; NAME/KEY: exon
; LOCATION: 25520..26016
; OTHER INFORMATION: exon8
US-09-218-207-183
                                                                                                                                                                                                                                                                                                                                                                                                   US-08-920-422-17
                                                                                                                                                                                                                                                                                                                                               Sequence 17, Application US/08920422A Patent No. 6255473
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.0%;
Matches 16; Conservative
                                                               Matches
                                                                                               Query Match
                                                                                                                                                                                                                  APPLICANT: Vitek, Michael P.
APPLICANT: Mitsuda, No. 6255473iaki
APPLICANT: ROSES, Allen D.
TITLE OF INVENTION: Presenilin-1 Gene Promoter
FILE REFERENCE: VITEKPRESENTLIN
CURRENT APPLICATION NUMBER: US/08/920,422A
CURRENT FILING DATE: 1997-08-29
NUMBER OF SEQ ID NOS: 22
                                                                                                                                                                                                        SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: exon
LOCATION: 5259..5328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: exon7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: exon
LOCATION: 23387..23510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: exon
LOCATION: 21789..21950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: exon
LOCATION: 14621..14710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: exon
LOCATION: 12675..12791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
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                              140 TGTCAGAAAGAAGTGA 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: exon6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: exon5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    VAME/KEY: exon
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: exon3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 142 TCAGAAAGAAGTGATG 157
                                                       Local Local 16;
                                                                            Similarity
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                                                               Conservative
                                                          4.3%; Score 16;
100.0%; Pred. No.
tive 0; Mismatc
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                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                            DB 4;
                                                            0;
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                                                                                        Length 48974;
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Qy 45 AAATCCAGTGCTACCA 60
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Db 42450 AAATCCAGTGCTACCA 42465
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                                                                                                                                                                                                            Sequence 57, Application US/09453702B Patent No. 6365723 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/09009913
Patent No. 6087485
GENERAL INFORMATION:
APPLICANT: AxyS Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM COMPUTER: COMPUTER: COMPUTER: DOS
SOTUMARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA: US/09/009,913
PTITING DATE: 21-JAN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Sherwood, Pamela J
REGISTRATION NUMBER: 36,677
REFERENCE/DOCKET NUMBER: SEY
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-327-3231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 72928 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Bozicevic & Reed, LLP
STREET: 285 Hamilton Ave, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION: NAME: Sherwood, Pamela J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS: dou
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
ZIP: 943
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 650-327-3231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Quarles & Brady
STREET: 1 South Pinckney Street
                                                                                               TITLE OF INVENTION: NO. 6365723el Sequences of E. coli 0157 NUMBER OF SEQUENCES: 265
                                                                                                                                                                                         APPLICANT: Blattner, Frederick R.
                                                                                                                                                                                                                                                                                                                                                                                               16;
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             CITY: Madison
STATE: WI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleic acid
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CA
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                                                                                                                                               Burland, Nicole T.
                                                                                                                     Pernu,
Plunkett, bu,
Ah, Rod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Genomic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                double
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ZIP: 53701-2113

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; TOPOLOGY: Inear;; MOLECULE TYPE: DNA (genomic); SEQUENCE DESCRIPTION: SEQ ID NO: 57: US-09-453-702B-57
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US-08-368-803-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 5733554
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 251-5000
TELEFAX: (608) 251-9166
INFORMATION FOR SEQ ID NO: 57:
                                                                                                                                                                                                                                                                                                                                                          APPLICANT: DUIMA:, C. APPLICANT: LAPLACE, Ellane L
APPLICANT: RIVIERE, Michel A
TITLE OF INVENTION: Avian Herpesvirus-based live recombinant avian
TITLE OF INVENTION: vaccine, in particular against Gumboro disease
TITLE OF INVENTION: vaccine, in particular against Gumboro disease
                             FILING DATE: 05-JAN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: SARRO, Thomas
                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         311 CCCAAGAAAAGGGGAA 326
                  TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                               STREET: 727 SOUTH 23RD STREET CITY: ARLINGTON
                                                                                               APPLICATION NUMBER: FILING DATE: 05-JA
                                                                                                                                                                                                                                                                              COUNTRY:
TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 60/110,955
FILING DATE: 04-DEC-1998
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US/09/453,702B
FILING DATE: 03-Dec-1999
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch. 1.44Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                             22202
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REGISTRATION NUMBER: 27386
REFERENCE/DOCKET NUMBER: 960296.95017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Word Perfect 8.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 87563
                                                                                                                                                                                                                                                                                                VIRGINIA
                                                                                                                                                                                                                                                                              USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DARTEIL, Raphael J
DUINAT, Carole V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BUBLOT, Michel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AUDONNET, Jean-Christophe F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                        LARSON AND TAYLOR
(703) 920-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.3%; Score 16; DB 4;
100.0%; Pred. No. 50;
                                                                                                                  US/08/368,803
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US-08-578-096A-22
                                                                                                                                                                            ; TOPOLOGY: linear; MOLECULE TYPE: other nucleic acid US-08-578-096A-22
Search completed: November 7, 2002, 23:12:13 Job time: 123.827 secs
                                                         Ъ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 22, Application US/08578096A Patent No. 5980906 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 36 base pairs
TYPE: nucleic acid
                                                                                                                                                 Query Match
                                                                                                                     Matches
                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Avian herpesylTITLE OF INVENTION: avian vaccine NUMBER OF SEQUENCES: 28 COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  194 GCATCATCAGCGGCC 208
                                                                                       194 GCATCATCAGCGGCC 208
                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
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les 15; Conserv
                                                                                                                                                                                                                                       TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/578,096A
                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                   Local
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                                                                                                                                                                                                                                                   LENGTH:
                                                           w
                                                         GCATCATCAGCGGCC 17
                                                                                                                     11 Similarity 100.
15; Conservative
                                                                                                                                                                                                                                                        36 base pairs
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100.0%;
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; Pred. No.
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Pred. No.
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                                                                                                                        Mismatches
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hes 0;
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1.6e+02;
                                                                                                                         0;
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                                                                                                                                                    Length 36;
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OM nucleic - nucleic search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Post-processing: Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Perfect score:
                                                                                                                                                                                                                                                                                                                                                                               Result
                                                                                                                            O
                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Published_Applications_NA:*

1: /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*

3: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*

5: /cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*

6: /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*

7: /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*

8: /cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*

9: /cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*

10: /cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*

11: /cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*

12: /cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*

13: /cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*

14: /cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*

14: /cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*
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US-09-825-294-214
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US-09-294-0938-5709
US-09-294-0938-5021
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-09-764-869	-09-764-877-	US-09-764-860-1156	-291-	US-09-922-261-359	US-09-802-669-94	US-09-822-849A-343	US-09-801-368-41	US-09-813-358-34	US-09-815-242-8625	US-09-815-242-4715	US-10-001-887-14	US-09-833-381-94	US-09-764-864-423	US-09-393-634-22	US-09-864-761-10902	US-09-983-965-3483	US-09-960-352-11005	US-09-960-352-3690	US-10-103-859-1	US-10-057-275-3	US-09-770-791-278	US-09-960-352-11486	us-09-960-352-1111	US-09-878-574-7775	US-09-923-876-3098
2170,	Sequence 2431, Ap	Sequence 1156, Ap	Sequence 1, Appli	Sequence 359, App	Sequence 94, Appl	Sequence 343, App	Sequence 41, Appl		1.5	4715,	14,	Sequence 94, Appl	423	Sequence 22, Appr	1090	Sequence 3483, Ap	-		Sequence 1, Appli	Sequence 3, Appli	Sequence 278, App	Sequence 11486, A		Sequence 7775, Ap	•

ALIGNMENTS

APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
APPLICANT: Filing, Steven P.
APPLICANT: Filing, Steven P.
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.4894C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 199
LENGTH: 369 ORGANISM: Homo sapiens
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FEATURE δÃ US-09-825-294-199 RESULT 1 Вþ 밁 δÃ В Sequence 199, Application US/09825294 Patent No. US20020004491A1 GENERAL INFORMATION: Query Match
Best Local Similarity 100.0%; Matches TYPE: DNA 121 GGTGAACGTTCAAGACATGTGTCAGAAAGAGTGATGGAGCAAAGTGCCGGGATCATGTA 180
121 GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA 180 369; Conservative 0; Score 369; DB 10; Pred. No. 4.5e-188; ); Mismatches 0; Length 369; Indels 0; Gaps

```
APPLICANT: X. Jiangchun

APPLICANT: Stolk, John A.

APPLICANT: Algate, Paul A.

APPLICANT: Algate, Paul A.

APPLICANT: Fling, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE

TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA

FILE REFERENCE: 210121.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FASTSEQ for Windows Version 3.0

SEQ ID NO 205

LENGTH: 1619
US-09-825-294-211
Sequence 211, Application US/09825294
Patent No. US20020004491A1
GENERAL INFORMATION:
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; ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                          GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT 269
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: XU, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTMARE: FastSEQ for Windows Version 3.0
SEQ ID NO 214
LENGTH: 1897
TYPE: DNA
TYPE: DNA
                                         US-09-825-294-214
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; Patent No. US20020004491A1
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-825-294-211
   Query Match
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APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C5
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 211
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Best Local
                                                       LOCATION: (1)...(1897)
OTHER INFORMATION: n =
                                                                          NAME/KEY: misc_feature LOCATION: (1)...(1897)
                                                                                                                ORGANISM: Homo sapiens FEATURE:
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GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fiing, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 215
SOFTWARE: F8stSEQ for Windows Version 3.0
SEQ ID NO 9
LENGTH: 396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9, Application US/09825294 Patent No. US20020004491A1
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                                                                                                                                                                                                                                                           Matches 281;
                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (1)...(396)
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
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                                 210
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                                                                                                                                                                                                                                                                          Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT 269
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTGCCTCGGCCCTCA 623
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                                                                                 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTTGTGCATCATCAGCGGCCT 209
                                                                                                                                            GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
                                                                  AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                                                              GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
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                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                        76.2%; score 281; DB 10; Length 3: 100.0%; Pred. No. 5.8e-141; vative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                         DB 10; Length 396;
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                         NUMBER OF SEQ ID NOS: 215
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 213
LENGTH: 480
TYPE: No. 2
                                                                                                GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION UMBER: US/09/825,294
CURRENT ETLING DATE: 2001-04-03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                               Sequence 213, Application US/09825294 Patent No. US20020004491A1
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        ORGANISM: Homo
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        sapiens
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APPLICANT: Algate, Paul A.

APPLICANT: Filing, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE

TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA

FILE REFERENCE: 210121.484C5

CURRENT APPLICATION UNUBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 212

LENGTH: 1010
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                                                                                                                                                                                                                                                                                                              318 CTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAGA 377
498 CATCAGCTGCTACACCCCCTCTTTGTAAC
                        271 CATCAGCTGCTGCAACACCCCTCTTTGTAAC 301
                                                                                                     438 TCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTG
                                                                                                                                 211 TCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTG
                                                                                                                                                                                                                                                             151 AGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTG 210
                                                                                                                                                                                                                                                                                                                                          91 CTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAGA 150
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US-09-294-093B-5287

; Sequence 5287, Application US/09294093B

; Patent No. US20010051335A1
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                                                                                                                          SOFTWARE:
SEQ ID NO 5287
                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/294,093B CURRENT FILING DATE: 1999-04-16 PRIOR APPLICATION NUMBER: 60/082,567 PRIOR FILING DATE: April 21, 1998 NUMBER OF SEQ ID NOS: 6207
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                                                                                                                                                                                                                                                          APPLICANT: Sherman, Bradley, K.
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN TASSEL
FILE REFERENCE: PL-0009 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: HATLOCKEY, SUSAN L.
APPLICANT: HATLOCKEY, SUSAN L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEO for Windows Version 4.0
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  NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No. US20010051335A1 700356180H1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Homo sapien
NAME/KEY: unsure
                                                         FEATURE:
                                                                       ORGANISM: Zea mays
                                                                                               TYPE: DNA
                                                                                                                ENGTH: 237
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 150 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCAT 197
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                                                                                                                                                  PERL Program
                                                                                                                                                                                                                                                                                                                                            Lalgudi, Raghunath, V.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14.9%; Score 55; DB 10;
100.0%; Pred. No. 5.6e-20;
tive 0; Mismatches 0;
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US-09-294-093B-5709
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; OTHER INFORMATION: a, t, c, g, or other
US-09-294-093B-5287
                                                                                  SOFTWARE:
SEQ ID NO 5021
LENGTH: 286
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                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/294,093B
CURRENT FILING DATE: 1999-04-16
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                                                                                                                                                                                                                          APPLICANT: Ito, Laura, Y.
APPLICANT: Sherman, Bradley, K.
TITLE_OF_INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN TASSEL
                                                                                                                                     PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: April
NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Sherman, Bradley, K.
TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN TASSEL
FILE REFERENCE: PL-0009 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: PRIOR FILING DATE: April
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/294,093B CURRENT FILING DATE: 1999-04-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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                                                  ORGANISM: Zea mays
                                                                        TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: unsure
LOCATION: 19, 21, 234, 245
OTHER INFORMATION: a, t, c, g, or other
OTHER INFORMATION: Incyte ID No. US20010051335A1 700355590H1
                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 256
TYPE: DNA
ORGANISM: Zea mays
                   NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Incyte ID No. US20010051335A1 700382209H1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    192 GTGCATCATCAGCGGCCT 209
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                                                                                                                                                                                                                                                                                                                                                 Application US/09294093B
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                                                                                                                                                                                                                                                                                           Lalgudi, Raghunath, V.
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                                                                                                                                                   April 21,
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100.0%; Pred. No.
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Pred. No.
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; LOCATION: 8, 26, 57, 117, 124, 216, 248,
; OTHER INFORMATION: a, t, c, g, or other
US-09-294-093B-5021
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SEQ ID NO 6
LENGTH: 1791
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Best Local
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APPLICANT:
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                      SOFTWARE: PatentIn Ver.
SEQ ID NO 682
LENGTH: 1298
TYPE: DNA
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CURRENT FILING DATE: 1999-10-12
CURRENT APPLICATION NUMBER: US/09/416,384A
PRIOR APPLICATION NUMBER: 60/106,457
PRIOR FILING DATE: 1999-10-30
PRIOR APPLICATION NUMBER: 60/103,955
PRIOR FILING DATE: 1998-10-12
PRIOR FILING DATE: 1998-10-12
                                                                                           CURRENT APPLICATION NUMBER: US/09/925,300
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05988
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEG ID NOS: 1890
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PRIOR FILING DATE: 1999-05-03
NUMBER OF SEQ ID NOS: 71
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TITLE OF INVENTION: Genes, proteins and biallelic markers related to central...
FILE REFERENCE: GENEE..045A0.
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APPLICANT: Steve Ruben
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA101
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                192 GTGCATCATCAGCGGCCT 209
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nes 18; Conserv
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Local Similarity 100.0%; Score 18; DB
Local Similarity 100.0%; Pred. No. 3.7
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Pred. No.
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; NAME/KEY: misc feature
; LCGATION: (1294)
; OTHER INFORMATION: n equals a,t,g, or c
US-09-925-300-682
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; LOCATION: (1)...(1713)
US-09-815-242-7853
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CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-10-23
PRIOR FILING DATE: 2000-10-23
                                                                                                  RESULT 15
US-09-815-242-6571
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APPLICANT: Haselbeck, Robert
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Patent No. US20020061569A1
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Best Local Similarity
Matches 17; Conserv
Sequence 6571, Application US/09815242
Patent No. US20020061569A1
GENERAL INFORMATION:
APPLICANT: Haselbeck, Robert
APPLICANT: Ohlsen, Karl L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Fas
SEQ ID NO 7853
                                                                                                                                                                                                                                                                                             Query Match
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TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
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NUMBER OF SEQ ID NOS: 14110
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PRIOR FILING DATE: 2000-11-27
                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 1713
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Xu, H. Howard
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Zyskind, Judith W.
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100.0%; Pred. No. 13;
tive 0; Mismatches
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Pred. No.
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. 13;
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; TYPE: DNA; ORGANISM: Homo sapiens US-09-964-824A-313
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; LOCATION: (1)...(2649)
US-09-815-242-6571
                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: HOURIGAN, Stephen
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Signatu
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/09/964,824A
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
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PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
PRIOR FILING DATE: 2000-09-28
         Matches
                                                                 Query Match
                                                                                                                                                                                                                           SEQ ID NO 313
LENGTH: 6252
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PRIOR FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
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PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR FILING DATE: 2000-05-26
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TITLE OF INVENTION: Identification of Essential Genes
TITLE OF INVENTION: Prokaryotes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/: PRIOR FILING DATE: 2001-02-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/815,242 CURRENT FILING DATE: 2001-03-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 2649
TYPE: DNA
ORGANISM: Enterococcus faecalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 14110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: ELITRA.011A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2563 AAAGAAGTGATGGAGCA 2579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 146 AAAGAAGTGATGGAGCA 162
Y Match 4.6%; Score 17; DB 10; Length 6252; Local Similarity 100.0%; Pred. No. 13; hes 17; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zyskind, Judith W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yamamoto, Robert T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Trawick,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Daniel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                John D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 60/269,308
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. 13;
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Gaps
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RESULT 18
US-09-764-860-1187
; Sequence 1187, Application US/09764860
; Patent No. US20020094953A1
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US-09-070-927A-121
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                                                                                   GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                Matches
                   APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC008
CURRENT APPLICATION NUMBER: US/09/764,860
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 121
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                            7402 AAAGAAGTGATGGAGCA 7418
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                                                                                                                                                                                                                                                                                                              Local Similarity
nes 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE DESCRIPTION: SEQ ID NO: 121:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Enterococcus faecialis Polynucleotides and Polypeptides NUMBER OF SEQUENCES: 982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Charles A. Kunsch
Patrick J. Dillon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/046,655
FILING DATE: 1997-05-16
APPLICATION NUMBER: 60/044,031
FILING DATE: 1997-05-06
APPLICATION NUMBER: 60/066,009
FILING DATE: 1997-11-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/070,927A FILING DATE: 04-May-2000 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Kenley K. Hoover
REGISTRATION NUMBER: 40,302
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OPERATING SYSTEM: MSDOS version 6.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 8033 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Human Genome Sciences, Inc
STREET: 9410 Key West Avenue
                                                                                                                                                                                                                                                                                                              Conservative
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100.0%; Pred. No.
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US-09-815-242-2587
; Sequence 2587, Application US/09815242
; Patent No. US20020061569A1
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               RESULT 20
US-09-923-876-3098/c
Sequence 3098, Application US/09923876
; Patent No. US20020013958A1
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                                                                                                                                                                                                                                                                                ; ORGANISM: Staphylococcus aureus US-09-815-242-2587
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SEQ ID NO 1187
LENGTH: 11842
TYPE: DNA
ORGANISM: Homo sapiens
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Best Local Similarity
Matches 17; Conserv
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SOFTWARE: FastSEQ for Windows Version 4.0
SEC ID NO 2587
LENGTH: 124
GENERAL INFORMATION
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CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR FILING DATE: 2000-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: 60/269,308
PRIOR APPLICATION NUMBER: 60/269,308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Xu, H. Howard
TITLE OF INVENTION: Identification of Essential Genes
TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
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                                                                                                                                                                                                                                                                                                                      TYPE: DNA
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                                                                                                                               41 AATCCAGTGCTACCAG 56
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nes 16; Conserv
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Trawick, John D.
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                                                                                                                                                                                                        Conservative
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100.0%; Pred. No.
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100.0%; Pred. No.
tive 0; Mismatc
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                                                                                                                                                                                                                                           Length 124;
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US-09-923-876-3098
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; OTHER INFORMATION: Clone ID: 701100324H1
US-09-878-574-7775
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                                                                                                                                                                                 US-09-960-352-1111
                                                                                                                                                                                                   RESULT 22
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APPLICANT: Thompson, Michael D.

TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with

TITLE OF INVENTION: Plants

FILE REFERENCE: 38-21(15401)B

CURRENT APPLICATION NUMBER: US/09/878,574

CURRENT FILING DATE: 2001-11-21

PRIOR APPLICATION NUMBER: 09/333,535

PRIOR APPLICATION NUMBER: 09/333,535

PRIOR FILING DATE: 1999-06-14

NUMBER OF SEQ ID NOS: 15775

SEQ ID NO 7775

LENGTH: 271
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SEQ ID NO 3098
LENGTH: 236
TYPE: DNA
GENERAL INFORMATION:
APPLICANT: Warren, Wesley C.
APPLICANT: Tao, Nengbing
APPLICANT: Syatt, John C.
APPLICANT: Mathialagan, Nagappan
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                  Sequence 1111, Application US/09960352 Patent No. US20020137139A1
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CURRENT FILING DATE: 2001-08-06
PRIOR APPLICATION NUMBER: 09/298,329
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APPLICANT: Sherman, Bradley K.
TITLE OF INVENTION: POLYUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN SEEDLING
FILE REFERENCE: PL-0012-1 CON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 1999-04-21
PRIOR APPLICATION NUMBER: 60/085,331
PRIOR FILING DATE: 1998-05-05
NUMBER OF SEQ ID NOS: 6332
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE: NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No. US20020013958A1 700161660H1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        131 TCAGCTGCTGCAACAC 116
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                                                                                                                                                                                                                                                                                                              228 GGTACCAGTCCTTCTG 243
                                                                                                                                                                                                                                                                    104 GGTACCAGTCCTTCTG 119
                                                                                                                                                                                                                                                                                                                                                           16;
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                    4.3%; Score 16; DB 10; 100.0%; Pred. No. 42;
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                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB
42;
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                                                                                                                                                                                                                                                                                                                                                                                                         Length 271;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 236;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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                                                                                                                                                                                                                                                                                                                                                               Gaps
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밁
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OTHER INFORMATION: Clone ID: 49-LIB34-018-Q1-E1-E10
US-09-960-352-11486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; LENGTH: 364
; TYPE: DNA
; ORGANISM: Bos taurus
; OTHER INFORMATION: Clone ID: 05-LIB34-056-Q1-E1-B1
US-09-960-352-1111
                                                                                                                                                                                                                                                                                                          US-09-770-791-278/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-960-352-11486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:

APPLICANT: Wairen, Wesley C.

APPLICANT: Wairen, Wesley C.

APPLICANT: Byatt, John C.

APPLICANT: Byatt, John C.

APPLICANT: Mathialagan, Nagappan

TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND

TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION

FILE REFERENCE: 16511.006/37-21(10298)C

CURRENT APPLICATION NUMBER: US/09/960,352

CURRENT FILING DATE: 2001-09-24

NUMBER OF SEQ ID NOS: 15112

SEQ ID NO 11486

LENGTH: 369
                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                          Sequence 278, Application US/09770791 Patent No. US20020062014A1
                                                                                                                                                                             APPLICANT: APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 11486, Application Patent No. US20020137139A1
                                                                                                                               APPLICANT:
APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 1111
LENGTH: 364
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                            APPLICANT:
                      APPLICANT:
                                                                APPLICANT:
                                                                                                APPLICANT:
                                                                                                                                                                                                                                             APPLICANT: Gorlach, Jorn
                                   APPLICANT:
                                                    APPLICANT:
                                                                                   APPLICANT:
                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION FILE REFERENCE: 16511.006/37-21(10298)C CURRENT APPLICATION NUMBER: US/09/960,352 CURRENT FILLING DATE: 2001-09-24 NUMBER OF SEQ ID NOS: 15112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 194 CTGCAAATCCAGTGCT 209
                                                                                                                                                                                                                                                                                                                                                                           41 CTGCAAATCCAGTGCT 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               41 CTGCAAATCCAGTGCT 56
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                                                                                          Matthew, Abraham V
Ledford, Brooke L.
                                                                                                                                                                                          An, Yong-Qiang
Hamilton, Carol M.
Price, Jennifer L.
               Slader,
                                                              Woessner, Jeffrey P.
Haas, William David
Davis, Keith R.
                            Kricker, Maja
                                             Garcia, Carlos A.
                                                                                                                                 Page,
                                                                                                                                            Yu, Yang
Rameaka, Joshua G.
                                                                                                                                                                             Raines, Tracy M.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                            Amy
                                                                                                           Abraham V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.3%;
100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 16;
Pred. No.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
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RESULT 25
US-10-057-275-3/c
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; ORGANISM: Arabidopsis thaliana
US-09-770-791-278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  В
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Patent No. US20020155545A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 278
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Best Local Similarity
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Hurban, Patrick
TITLE OF INVENTION: Expressed Sequences of Arabidopsis
TITLE OF INVENTION: thaliana
FILE REFERENCE: 2029 (PARA-018PRV)
CURRENT APPLICATION NUMBER: US/09/770,791
CURRENT FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: 60/178,480
PRIOR APPLICATION NUMBER: 60/178,480
PRIOR FILING DATE: 2000-01-27
NUMBER OF TIME DATE: 2000-01-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 999
                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 375
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      80 AACAACGACTGCTCCT 95
                                                                                                                                                                                                                               NAME: Luther, Barbara J.
REGISTRATION NUMBER: 33,954
REFERENCE/DOCKET NUMBER: PF-0027 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-852-0195
                  MOLECULE TYPE: cDNA
IMMEDIATE SOURCE:
LIBRARY: Human Pancreas
CLONE: 226152
SEQUENCE DESCRIPTION: SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: DOS
SOFTWARE: PRASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/10/057, 275
FILING DATE: 25-Jan-2002
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc
STREET: 3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: NEW CHEMOKINES EXPRESSED IN PANCREAS NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Coleman, Roger
Bandman, Olga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16;
                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/390,740A FILING DATE: February 17, 1995
                                                                                                       TOPOLOGY: linear
                                                                                                                                                                         LENGTH: 402 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hoffman,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Allen, Keith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.3%; Score 16; ilarity 100.0%; Pred. No. Conservative 0; Mismatcl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      U.S.
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APPLICANT: Byatt, John C.

APPLICANT: Mathhalagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
FILE REFERENCE: 16511.006/37-21(10298)C
CURRENT APPLICATION NUMBER: US/99/960,352
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 15112
SEQ ID NOS: 15112
LENGTH: 406
TYPE: n...
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; LOCATION: (1)..(405)
; OTHER INFORMATION:
US-10-103-859-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/10103859 Patent No. US20020155094A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/103,859
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: US 09/496,273
PRIOR FILING DATE: 2000-02-01
PRIOR APPLICATION NUMBER: US 08/852,212
PRIOR FILING DATE: 1997-05-06
PRIOR FILING DATE: 1996-05-14
PRIOR FILING DATE: 1996-05-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Pelus, L.M.
TITLE OF INVENTION: Method of Treating Sepsis and ARDS Using Chemokine Beta-9
FILE REFERENCE: 1488.1520003
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                 TYPE: DNA
ORGANISM: BOS taurus
OTHER INFORMATION: Clone ID: 16-LIB34-030-Q1-E1-D11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                263 TGCATCAGCTGCTGCA 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 269 TGCATCAGCTGCTGCA 284
                                                                                                                                                                                                                                                                                                                                                                                                                                                    263 TGCATCAGCTGCTGCA 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     269 TGCATCAGCTGCTGCA 284
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4.3%; Score 16; DB 9;
100.0%; Pred No. 43;
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100.0%;
 4.3%;
100.0%;
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 DB 10;
43;
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                  Length 406;
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LOCATION: (333),(347),(373),(384)
; OTHER INFORMATION: unsure at all n locations
; OTHER INFORMATION: Clone ID: 47-LIB34-011-Q1-E1-D4
US-09-960-352-11005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-960-352-11005
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LENGTH: 411
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Byatt, John C. APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILE REPERENCE: 16511.006/37-21(10298)C
CURRENT APPLICATION NUMBER: US/09/960,352
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 15112
                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 5912 SEQ ID NO 3483
Query Match
                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Warren, Wesley C.
APPLICANT: Tao, Nengbing
APPLICANT: Byatt, John C.
APPLICANT: Byatt, John C.
APPLICANT: Wathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILE REFERENCE: 37-21(10297)C
CURRENT APPLICATION NUMBER: US/09/983,965
CURRENT FILING DATE: 201-10-26
CURRENT FILING DATE: 201-10-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Warren, Wesley C. APPLICANT: Tao, Nengbing
                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 09/465,231
PRIOR FILING DATE: 1999-12-15
PRIOR APPLICATION NUMBER: US 60/113,678
PRIOR FILING DATE: 1998-12-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
                                                              FEATURE:
NAME/KEY: unsure
LOCATION: (76),(81),(87)
OTHER INFORMATION: unsure at all n locations
OTHER INFORMATION: Clone ID: 37-LIB3058-007-Q1-K1-B2
                                                                                                                                                                            ORGANISM: Bos taurus
                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                        ENGTH: 414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           205 CTGCAAATCCAGTGCT 220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          389 CTGCAAATCCAGTGCT 404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     41 CTGCAAATCCAGTGCT 56
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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100.0%; Pred. No.
    4.38;
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    Score 16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 10; Length 411; 43;
    DB 10; Length 414;
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; OTHER INFORMATION: MAP TO ACO03103.1
; OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1;
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 0.9
; OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 0.89
; OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.2
; OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.6
; OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.1
; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-864-761-10902/c
                                                                                                                                                                                                                                                                                               SOFTWARE: Annomax Sequence Listing Engine vers. SEQ ID NO 10902 LENGTH: 488
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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Patent No. US20020048763A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY FILE REFERENCE: Acomica-x-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 09/632,366 PRIOR FILING DATE: 2000-08-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                       ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/180,312
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FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 09/608,408 FILING DATE: 2000-06-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 2000-09-21
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APPLICATION NUMBER: PCT/US01/00662
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT,
FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/US01/00665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: PCT/US01/00667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/US01/00666 FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 60/236,359 FILING DATE: 2000-09-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: GB 24263.6 FILING DATE: 2000-10-04
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Chen, Wensheng
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RESULT 32
US-09-764-864-423/c
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APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and
FILE REFERENCE: PTZ23
                                                                                                                                                                                                                        SOFTWARE: PatentIn Ver. SEQ ID NO 423
                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 423, Application US/09764864 Patent No. US20020132753A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 16; Conserv
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SEQ ID NO 22
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                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/764,864
CURRENT FILING DATE: 2001-01-17
                                                                                                                                                                                                                                                                 Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 1792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: The Regents of the University of California APPLICANT: The Government of the United States of Amer: APPLICANT: as represented by the Secretary of the APPLICANT: Department of Health and Human Services TITLE OF INVENTION: SF, a No. US20020051997Alel Family FILE REFERENCE: 02307E-098000US
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Best Local Similarity
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CURRENT FILING DATE: 1999-09-10
NUMBER OF SEQ ID NOS: 92
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                                                                                       LOCATION: (513)
OTHER INFORMATION: n equals a,t,g,
                                                                                                                                                                                           TYPE: DNA
NAME/KEY: SITE LOCATION: (569)
                                   OTHER INFORMATION: n equals a,t,g, or
                                                        NAME/KEY: SITE
LOCATION: (536)
                                                                                                                                FEATURE:
NAME/KEY: SITE
                                                                                                                                                                      ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Mus sp.
FEATURE:
                                                                                                                                                                                                             ENGTH: 615
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             138 TGTGTCAGAAAGAAGT 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            175 TGTGTCAGAAAGAAGT 190
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                                                          (536)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PatentIn Ver. 2.1
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Ryba, Nick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hoon, Mark
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100.0%;
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%; Pred. No. 43;
0; Mismatches
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 10; Length 538;
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US-09-833-381-94
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APPLICANT: Robison, Keith E.
APPLICANT: Robison, Keith E.
TITLE OF INVENTION: No. US20020132090Alel Nucleic Acid and Protein Homologs
FILE REFERENCE: 5800-119
CURRENT APPLICATION NUMBER: US/09/833,381
CURRENT FILING DATE: 2001-04-11
PRIOR APPLICATION NUMBER: 09/516,448
PRIOR FILING DATE: 2000-02-29
                                                                                                                                                                                                            APPLICANT: Salceda, Susana
APPLICANT: Macina, Roberto
APPLICANT: Recipon, Herve
APPLICANT: Cafferkey, Robert
APPLICANT: Sun, Yongming
APPLICANT: Liu, Chenghua
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 2050
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 94
LENGTH: 654
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 94, Application US/09833381 Patent No. US20020132090A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches 16;
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                                                                                                                                                                                                                                                                                                                                       Sequence 14, Application US/10001887 Patent No. US20020155464A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 4.3%; Score 16; Best Local Similarity 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                            APPLICANT: Sun, Yongming
APPLICANT: Liu, Chenghua
TITLE OF INVENTION: Compositions and Methods Relating to Breast Specific Genes and
EILE REFERENCE: DEX-0269
EILE REFERENCE: DEX-0269
CURRENT FILING DATE: 2001-11-20
PRIOR APPLICATION NUMBER: 60/249,998
PRIOR FILING DATE: 2000-11-20
PRIOR APPLICATION NUMBER: 60/252,563
PRIOR FILING DATE: 2000-11-22
NUMBER OF SEQ ID NOS: 137
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: SITE LOCATION: (588)
OTHER INFORMATION: n
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LOCATION: (610)
OTHER INFORMATION: n equals a,t,g, or
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LOCATION: (597)
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LENGTH: 689
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                  NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16;
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100.0%;
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Pred. No.
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43;
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 APPLICANT:
APPLICANT:
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RESULT 35
US-09-815-242-4715/c
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CURRENT EILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR FILING DATE: 2000-03-23
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/203,625
PRIOR FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR APPLICATION NUMBER: 60/253,931
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2000-12-22
                                                                                                                                           RESULT 36
US-09-815-242-8625/c
                                                                                                                                                                                                                                                                                                                                                                               ; ORGANISM: Staphylococcus aureus US-09-815-242-4715
                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 4715
LENGTH: 1172
TYPE: DNA
Sequence 8625, Application US/09815242
Patent No. US20020061569A1
GENERAL INFORMATION:
APPLICANT: Haselbeck, Robert
APPLICANT: Ohlsen, Kari L.
APPLICANT: Zyskind, Judith W.
APPLICANT: Wall, Daniel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Yamamoto, Robert T.
APPLICANT: Xu, H. Howard
TITLE OF INVENTION: Identification of Essential Genes in
TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 14110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/269,308
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nes 16; Conservative
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[00.0%; Pred. No.
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Trawick, John D. Carr, Grant J.

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; LOCATION: (1)...(1179)
US-09-815-242-8625
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                                                                                      Matches
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 34
LENGTH: 1549
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Best Local Similarity 100.0%; F
Matches 16; Conservative 0;
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PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR FILING DATE: 2000-05-26
                                                                                                                Best
                                                                                                                                  Query Match
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SEQ ID NO 8625
LENGTH: 1179
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APPLICANT: Pyle, Ruth
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN AND ENDOMETRIAL CANCER
FILE REFERENCE: 210121-501
CURRENT APPLICATION NUMBER: US/09/813,358
CURRENT FILING DATE: 2001-03-21
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PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2000-12-22
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APPLICANT: Xu, H. Howard
TITLE OF INVENTION: Identification of Essential Genes in
TITLE OF INVENTION: Prokaryotes
FILE REFERENCE: ELITRA.011A
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CURRENT FILING DATE: 2001-03-21
                                                                                                                                                                                         FEATURE: MISC_feature LOCATION: (1)...(1549)
OTHER INFORMATION: n - A,T,C or G
                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                         TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Staphylococcus aureus FEATURE:
                                         277 CTGCTGCAACACCCCT 292
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  57
                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     46 AATCCAGTGCTACCAG 61
CTGCTGCAACACCCCT 72
                                                                                      16;
                                                                                    Conservative
                                                                                                         4.3%;
                                                                                    0;
                                                                                                       Score 16;
Pred. No.
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                                                                                  Mismatches
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                                                                                                                           DB 10; Length 1549;
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CURRENT APPLICATION NUMBER: US/09/801,368
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: US 09/487,558
PRIOR FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: US 60/160,587
PRIOR APPLICATION NUMBER: US 60/160,587
PRIOR FILING DATE: 1999-10-20
NUMBER OF SEQ ID NOS: 440
SOFTWARE: Patentin version 3.0
SEQ ID NO 41
LENGTH: 1989
TYPE: DNA
ORGANISM: Aspergillus nidulans
US-09-801-368-41
US-09-822-849A-343
                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-822-849A-343/c
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                                                                    CURRENT APPLICATION NUMBER: US/09/82,849A CURRENT FILING DATE: 2001-09-04 PRIOR APPLICATION NUMBER: 60/195/582 PRIOR FILING DATE: 2000-04-06 NUMBER OF SEQ ID NOS: 598 SOFTMARE: Patentin Ver. 2.0 SEQ ID NO 343
                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. US20020045170A1
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 343, Application US/09822849A
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Best Local
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APPLICANT:
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Wong, Gordon G. APPLICANT: Clark, Hilary APPLICANT: Fechtel, Kim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Busby, APPLICANT: Cali, APPLICANT: Hecht
                                                                                                                                                                                                    APPLICANT: Genetics Institute, Inc.
TITLE OF INVENTION: POLYNUCLEOTIDES ENCODING NOVEL SECRETED PROTEINS
FILE REFERENCE: GIN 6403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Summers, Eric TITLE OF INVENTION: Methods for Improving Secondary Metabolite Production in Fungi-
FILE REFERENCE: 109272.147
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                 ORGANISM: Homo sapiens
                                    TYPE: DNA
                                                      LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1026 ACCGCAAGTCCTGTGC 1041
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Graham, James R.
                                                                                                                                                                                                                                                                                               Resnick, Richard J.
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Silva, Jeff
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Salama, Sofie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Milne, Todd
No. US20020128250Alman, Thea
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Maxon, Mary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cali, Brian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Holtzman, Doug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.3%;
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RESULT 40
US-09-802-669-94/c
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US-09-802-669-94
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-922-261-359
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 94, Application US/09802669 Patent No. US20020004490A1
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APPLICANT: Zhang, Hong
TITLE OF INVENTION: Antisense Compound Modulation of Fas Mediated Signaling
FILE REFERENCE: ISPH-545
CURRENT APPLICATION NUMBER: US/09/802,669
CURRENT FILING DATE: 2001-03-09
PRIOR APPLICATION NUMBER: US 09/665,615
PRIOR APPLICATION NUMBER: US 09/665,615
PRIOR APPLICATION NUMBER: US 09/290,640
PRIOR FILING DATE: 2000-09-18
PRIOR FILING DATE: 1999-04-12
NUMBER OF SEQ ID NOS: 180
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 359, Application US/09922261 Patent No. US20020111471A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                          SEQ ID NO 359
                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                           APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Dean, Nicholas M. APPLICANT: Marcusson, Eric G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE:
                                                                                          PRIOR APPLICATION NUMBER: US/09/461,697
PRIOR FILING DATE: 1999-12-14
NUMBER OF SEQ ID NOS: 466
SOFTWARE: EastSEQ for Windows Version 4.0
                                                                                                                                                                                                          APPLICANT: Portbury, Stuart D.
APPLICANT: Puranam, Kasturi
APPLICANT: Katz, Lawrence C.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING
TITLE OF INVENTION: CELL DEATH
TITLE OF INVENTION: CELL DEATH
FILE REFERENCE: 10001-005-999
                                                                                                                                                                     CURRENT FILING DATE: 2001-08-03
                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: COGENT NEUROSCIENCE, Inc. APPLICANT: Lo, Donald C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 2165
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: CDS
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                                                        LENGTH: 2702
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               306 CAAGGCCCAAGAAAAG 321
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                                                                                                                                                                                                                                                                                                                                               Barney, Shawn
Thomas, Mary Beth
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Conservative (
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US-09-834-291-1/c
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                                                                                                                                           ; TYPE: DNA; ORGANISM: Homo sapiens
US-09-764-860-1156
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
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Best Local Similarity
                                                                                                                                                                                           Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 1198
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 1156
LENGTH: 3294
                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1156, Application US/09764860 Patent No. US20020094953A1
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                                                                      Matches
                                                                                        Best Local Similarity
                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 1999-10-18
PRIOR APPLICATION NUMBER: DE 198 47 779.1
PRIOR FILING DATE: 1998-10-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/834,291
CURRENT FILING DATE: 2001-08-21
PRIOR APPLICATION NUMBER: PCT/DE99/03343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/764,860 CURRENT FILING DATE: 2001-01-17
                                                                                                                                                                                                                                                                                                                          APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PC008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 3212
TYPE: DNA
ORGANISM: Homo Sapiens
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819 GGTACCAGTCCTTCTG 834
                                  228 GGTACCAGTCCTTCTG 243
                                                                        16;
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100.0%; Pred. No. 44;
tive 0; Mismatches (
                                                                    4.3%; Score 16; DB
100.0%; Pred. No. 44;
tive 0; Mismatches
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                                                                                                           DB 10; Length 3294;
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RESULT 44 US-09-764-877-2431

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GENERAL INFORMATION:

APPLICANT: Rosen et al.

TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies

FILE REFERENCE: PC005

CURRENT APPLICATION NUMBER: US/09/764,877

CURRENT FILING DATE: 2001-01-17

Prior application data removed - refer to PALM or file wrapper NUMBER OF SEQ ID NOS: 4031

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 2431

LENGTH: 11749

TYPE: DNA
ORGANISM: Homo sapiens
US-09-764-877-2431
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 45
US-09-764-869-2170
; Sequence 2170, Application US/09764869
; Peatent No. US20020061521A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PC007
; FILE REFERENCE: PC007
                                                                                                                                    δδ
                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/764,869

CURRENT FILING DATE: 2001-01-17

Prior application data removed - refer to PALM or file wrapper NUMBER OF SEQ ID NOS: 2442

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 2170

LENGTH: 17581

TYPE: DNA

ORGANISM: HOMO Sapiens

US-09-764-869-2170
Search completed: November 8, 2002, 02:05:12 Job time: 41.0364 secs
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                                                                                                                                                                        Query Match 4.3%; Score 16; DB 10; Length 17581; Best Local Similarity 100.0%; Pred. No. 46; Matches 16; Conservative 0; Mismatches 0; Indels 0
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                                                                                      2661 AAGAAAAGGGGAAGTT 2676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                314 AAGAAAAGGGGAAGTT 329
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OM nucleic - nucleic search, using sw model
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Maximum DB seq length: 2000000000
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Perfect score:
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369
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        "Cgn2_6/ptodata/1/pna/US081_COMB.seq: *
(Cgn2_6/ptodata/1/pna/US081_COMB.seq: *
(Cgn2_6/ptodata/1/pna/US081_COMB.seq: *
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Copyright (c) 1993 - 2002 Compugen Ltd
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/cgn2_6/ptodata/1/pna/US06_COMB.seq:*
/cgn2_6/ptodata/1/pna/US07_COMB.seq:*
'cgn2_6/ptodata/1/pna/US102B_COMB
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/cgn2_6/ptodata/1/pna/US6001_COMB.seq:
/cgn2_6/ptodata/1/pna/US6002_COMB.seq:
/cgn2_6/ptodata/1/pna/US6003_COMB.seq:
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/cgn2_6/ptodata/1/pna/US6006_COMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.

Score

Match Length DB

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Sequence 15, Appl	Sequence 15, Appl	Sequence 15, Appl	Sequence 15, Appl	Sequence 15, Appl	Sequence 15, Appl	Sequence 15, Appl	Sequence 15, Appl		Sequence 18421, A	Sequence 6100, Ap	Sequence 18421, A	Sequence 283, App	•	•	•	Sequence 199, App	Sequence 199, App	Sequence 199, App	Sequence 199, App	Sequence 199, App	Description

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Sequence 199, Application PC/TUS0145395

GENERAL INFORMATION:
APPLICANT: CORIAG COrporation
APPLICANT: Xu, Jiangchun
APPLICANT: Xtolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Fling, Steven P.
APPLICANT: Fling, Steven P.
APPLICANT: Molesh, David Alan
APPLICANT: MOLESH, David Alan
APPLICANT: MOLESH DAVID AND METHODS FOR THE THE
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THE
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 21012.48401.PC
                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: PCT/US01/45395
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
LENGTH: 369
                                                                                      US-10-216-167-15
US-10-218-612-15
US-10-218-612-15
US-10-218-765-15
US-10-218-784-15
US-10-218-930-15
US-10-218-930-15
US-10-219-906-15
US-10-219-908-15
US-10-219-908-15
US-10-219-908-15
US-10-219-908-15
US-10-219-908-15
US-10-219-908-15
US-10-219-907-15
                                                                                                                             Score 369; DB 1; Pred. No. 1.1e-188; 0; Mismatches 0;
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                                                                                                                                              Length 369;
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APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

TITLE OF INVENTION: OVARIAN TUMOR SEQUENCES
TITLE OF INVENTION: METHODS OF USE THEREFOR
FILE REFERENCE: 210121.484C3
CURRENT FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 199
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 199
LENGTH: 369
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(369)
OTHER INFORMATION: n = A,T,C or G
US-09-656-668-199
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Pred. No. 1.1e-188;
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; FEATURE:
, NAME/KEY: misc\_feature
; LOCATION: 29, 345
; OTHER INFORMATION: n =
PCT-US01-45395-199

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Query Match Best Local 9 Matches

369;

Similarity

100.0%; ilarity 100.0%; Conservative

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SEQ

LENGTH: 369
TYPE: DNA
ORGANISM: Homo sapiens

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Sequence 199, Application US/09825294

GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun

APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

APPLICANT: Filing, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER FILE REFERENCE: 210121.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 199

LENGTH: 369
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APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CAN
FILE REFERENCE: 210121.484C4
CURRENT APPLICATION NUMBER: US/09/713,550
CURRENT FILING DATE: 2000-11-14
NUMBER OF SEQ ID NOS: 205
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 199
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
FEATURE: 569
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; LOCATION: (1)...(369)
; OTHER INFORMATION: n = A,T,C or
US-09-713-550-199
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US-09-825-294-199
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Pred. No. 1.1e-188;
; Mismatches 0;
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APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THI
TITLE OF INVENTION: COMPOSITIONS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C6
CURRENT APPLICATION NUMBER: US/09/970,966
CURRENT FILING DATE: 2001-10-02
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
LENGTH: 369
TYPE: DNA
ORGANISM: Homo sapiens
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US-09-970-966-199
Sequence 199, Application US/09970966
; GENERAL INFORMATION:
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FEATURE:
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NAME/KEY: misc_feature
LOCATION: 29, 345
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Fling, Steven P.
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llarity 100.0%;
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Pred. No. 1.1e-188;
); Mismatches 0;
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Pred. No. 1.1e-188;
Mismatches 0;
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; LOCATION: 29, 345
; OTHER INFORMATION: n =
US-10-212-677-199
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US-10-212-677-199
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Matches
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CURRENT FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 288
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 199
LENGTH: 369
                                                                                                                                                                                                                                                                                                          Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Fanger, Gary R.
APPLICANT: Harlocker, Susan L.
APPLICANT: McNeill, Patricia D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Chenault, Ruth A. APPLICANT: Xu, Jiangchun
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; OTHER INFORMATION: n e
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                                                                                                                                                                                                                                                                                                    Best Local Similarity Matches 317; Conserv
                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
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PRIOR FILING DATE: 2000-01-31
PRIOR APPLICATION NUMBER: 60/180,628
PRIOR ETLING DATE: 2000-02-04
NUMBER OF SEQ ID NOS: 1440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: PM002
CURRENT APPLICATION NUMBER: US/09/757,034
CURRENT FILING DATE: 2001-01-09
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NAME/KEY: SITE
LOCATION: (435)
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                                               AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
             CTGCCTCGGCCCTCANG 346
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RESULT 8 US-10-235-954-340

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PRIOR FILING DATE: 2001-01-09
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                                                                                                                                                                                                                                                      Sequence 283, Application US/10212677 GENERAL INFORMATION:
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CURRENT FILING DATE: 2002-09-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PM002C1N
APPLICANT: Chenault, Ruth A.
APPLICANT: Xu, Jiangchun
APPLICANT: Fanger, Gary R.
APPLICANT: Harlocker, Susan L.
APPLICANT: MCNEIll, Patricia D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature
LOCATION: (367)
OTHER INFORMATION: n equals
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OTHER INFORMATION: n equals
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100.0%;
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 Mismatches

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; TYPE: DNA; ORGANISM: Homo sapiens US-10-212-677-283
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US-09-234-611-18421
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LENGTH: 414
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TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-757
CURRENT APPLICATION NUMBER: US/09/234,611
CURRENT FILING DATE: 1999-01-21
NUMBER OF SEQ ID NOS: 21025
SOFTWARE: FASLSEQ for Windows Version 3.0
SEQ ID NO 18421
LENGTH: 488
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CURRENT APPLICATION NUMBER: US/10/212,677
CURRENT FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 288
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                    Matches 315;
                                                                                                                                              Query Match
                                                                                                                                                                                          ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(488)
OTHER INFORMATION: n = A,T,C or
                                                                                                                                                                                                                                                                    TYPE: DNA
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GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
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Similarity 100.0%;
15; Conservative (
                                                                                                                    Conservative
                                                                                                                                85.4%;
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; Pred. No. 2e-159;
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                                                                                                                    score 315; DB 16;
; pred. No. 2e-159;
0; Mismatches 0;
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RESULT 12
US-09-904-809-18421
Sequence 18421, Application US/09904809
GENERAL INFORMATION:
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: LOCATION: (1)...(488)
: OTHER INFORMATION: n = A,T,C
US-09-277-227-6100
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US-09-277-227-6100
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TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-766
CURRENT APPLICATION NUMBER: US/09/277,227
CURRENT FILING DATE: 1999-03-25
NUMBER OF SEQ ID NOS: 23600
SOFTWARE: FEATSEQ for Windows Version 3.0
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TYPE: DNA
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100.0%; Pred. No.
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; NAME/KEY: misc_feature
; LCCATION: (1)...(488)
; OTHER INFORMATION: n = A,T,C or
US-09-909-627-6100
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; GENERAL INFORMATION:
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                                                                                                                                                                  APPLICANT: Hyseq, Inc.

TITLE OF INVENTION: NOVEL NUCLEIC ACID SEQUENCES OBTAINED TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES FILE REFERENCE: 20411-766
CURRENT APPLICATION NUMBER: US/09/909,627
CURRENT FILING DATE: 2001-07-19
PRIOR APPLICATION NUMBER: 09/277,227
PRIOR FILING DATE: 1999-03-23
NUMBER OF SEQ ID NOS: 23680
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 6100
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CURRENT FILING DATE: 2001-07-12
PRIOR APPLICATION NUMBER: 09/234,611
PRIOR FILING DATE: 1999-01-22
NUMBER OF SEQ ID NOS: 21025
SOFTWARE: FBALSEQ for Windows Version 3.0
SEQ ID NO 18421
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Best Local
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                                                                                        FEATURE:
                                                                                                           ORGANISM: Homo sapiens
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TITLE OF INVENTION: HOVEL NUCLEIC ACID SEQUENCES OBTAINED
TITLE OF INVENTION: FROM VARIOUS CDNA LIBRARIES
FILE REFERENCE: 20411-757
                                                                                                                                        TYPE: DNA
                                                                                                                                                             LENGTH: 488
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TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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OTHER INFORMATION: n = A,T,C or G
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LENGTH: 1524
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                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/10/119,480
CURRENT FILING DATE: 2002-04-09
NUMBER OF SEQ ID NOS: 246
Prior Application removed - See File Wrapper or Palm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: P3530P1C1
                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
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                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Homo Sapien
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                                                                                                                                                    Local Similarity
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                                                 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
                                                                                           GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
                                 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
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Grimaldi, J. Christopher
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; SEQ ID NO 15
; LENGTH: 1524
; TYPE: DNA
; ORGANISM: HOMO Sapien
US-10-216-159A-15
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CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/06287
PRIOR APPLICATION NUMBER: 60/06287
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/06349
PRIOR APPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1998-03-20
                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/079728
PRIOR FILING DATE: 1998-03-27
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 246
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150 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
                                                                                                                                                 30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 89
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                                                                                                                                                                                                                                       Local Similarity
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                                                               GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAAG 149
                                        GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG
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Grimaldi, J. Christopher
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Gerritsen, Mary
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100.0%; Pred. No. 2e-159;
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US-10-216-160-15
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Best Local Similarity 100.
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APPLICANT: Desnoyers, Luc
APPLICANT: Gerritsen, Mary
APPLICANT: Goddard, Audrey
APPLICANT: Goddwski, Paul J.
APPLICANT: Grimaldi, J. Chri
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CURRENT APPLICATION NUMBER: US/10/216,160
CURRENT FILING DATE: 2002-08-09
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CTGCCTCGGCCCTCA 484
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                                                                     GCATCAGCTGCTGCAACACCCCCTCTTTGTAACGGGCCCAAGGCCCCAAGAAAAGGGGGAAGTT 329
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
TYPE: DNA
ORGANISM: Homo Sapien
US-10-216-162-15
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PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/06349
PRIOR APPLICATION NUMBER: 60/06349
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/064873
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/079910
PRIOR APPLICATION NUMBER: 60/079294
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PRIOR FILING DATE: 1998-03-26
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Best Local Similarity
Matches 315; Conserv
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APPLICANT:
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APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530P1C2
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CURRENT FILING DATE: 2002-08-09
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Goddard, Audrey
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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; ORGANISM: Homo Sapien
US-10-216-163-15
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CURRENT EILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/06287
PRIOR APPLICATION NUMBER: 60/06287
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
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FILING DATE: 1998-03-25
APPLICATION NUMBER: 60/079656
FILING DATE: 1998-03-26
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FILING DATE: 1997-10-31
                                     GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT 269
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FILING DATE: 1997-12-17
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Grimaldi, J. Christopher
Gurney, Austin L.
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Smith, Victoria
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CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/06287
PRIOR APPLICATION NUMBER: 60/06287
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
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LENGTH: 1524
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PRIOR FILING DATE: 1997-12-17
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PRIOR FILING DATE: 1998-03-20
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210 GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT 269
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                                                                                                                                                                           410
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                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/079294
FILING DATE: 1998-03-25
APPLICATION NUMBER: 60/079656
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 1998-03-26
                                   AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 349
                                                        AAGTGATGAAGCAAAGTGCCGGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
                                                                                                                                                                                                                                                             315;
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                                                                                                                                                                                                                                                                              Similarity
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Smith, Victoria
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Goddard, Audrey
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                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACIDS ENCODING THE SAME
                                                                                                                                                                                                                                                   85.4%; Score 315; DB 42; Length 1524; 100.0%; Pred. No. 2e-159; tive 0; Mismatches 0; Indels 0
                                                                                                                                                                                                                                                                                                                                                                                                                                   See File Wrapper or PALM
                                                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                 229
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150 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
        20
         90 GCTCCTCCCCGAGTTCATTGTGAATTGCACGG
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|GACT 229
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             AAG 149
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T 269
                                                                      Gaps
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	Qy 90 GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGACGGTCAAGACATGAAAAA 289 Db 230 GCTCCTCCCCCGGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTCAGAAAA 289
Dest Local Similarity 100.0%; Matches 315; Conservative  Qy 30 CAGGCTTTGCGCTGCAAATCCAG	30 170
; TYPE: DAM ; TYPE: DAM ; ORGANISM: Homo Sapien 0; US-10-216-166-15 85.	Ω
* Z P Y	; SEQ ID NO 15 ; LENGTH: 1524 ; TYPE: DNA ; ORGANISM: Homo Sapien US-10-216-165-15
PRIOR APPLICATION NUMBER: 60/079294 PRIOR FILLING DATE: 1998-03-25 PRIOR APPLICATION NUMBER: 60/079656 PRIOR FILLING DATE: 1998-03-26 PRIOR APPLICATION NUMBER: 60/079728	; PRIOR FILING DATE: 1998-03-26; PRIOR APPLICATION NUMBER: 60/079728; PRIOR FILING DATE: 1998-03-27; Remaining Prior Application data removed - See File Wrapper or PALM.; NUMBER OF SEQ ID NOS: 246
	; PRIOR FILING DATE: 1998-03-20 ; PRIOR APPLICATION NUMBER: 60/079294 ; PRIOR FILING DATE: 1998-03-25 ; PRIOR APPLICATION NUMBER: 60/079656
	; PRIOR FILING DATE: 1997-10-31 ; PRIOR APPLICATION NUMBER: 60/069873 ; PRIOR FILING DATE: 1997-12-17 ; PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: PRIOR APPLICATION NUMBER: PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1997-10	
CURRENT APPLICATION NUMBER: US/10/216,166 CURRENT FILING DATE: 2002-08-09 PRIOR APPLICATION NUMBER: 10/119,480 PRIOR FILING DATE: 2002-04-09 PRIOR FILING DATE: 2002-04-09	4-0 60 9-1
APPLICANT: WOOD, WILLIAM I. TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE PC TITLE OF INVENTION: ACIDS ENCODING THE SAME	; FILE REFERENCE: 93530PLC7; FILE REFERENCE: 93530PLC7; CURRENT APPLICATION NUMBER: US/10/216,165; CURRENT FILING DATE: 2002-08-09; CURRENT FILING DATE: 2002-08-09
; APPLICANT: Gurney, Austr ; APPLICANT: Smith, Victor ; APPLICANT: Stephan, Jean ; APPLICANT: Watanabe, Col	S I
APPLICANT: Goddard, Audrey APPLICANT: Godowski, Paul J. APPLICANT: Grimaldi, J. Christopher	APPLICANT: Suith, Victoria  APPLICANT: Smith, Victoria  APPLICANT: Stephan, Jean-Philippe F.  APPLICANT: Watanahe, Colin L.
HILL HILL HILL HILL HILL HILL HILL HILL	
RESULT 21 US-10-216-166-15 ; Sequence 15, Application US/10216166	_ = =
Oy 330 CTGCCTCGGCCCTCA 344	SULT 20 -10-216-165-1
QY 270 GCATCAGCTGCTACACCCCCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGAAATT	330 470
Qy 210 GTCTCATCGCCTCTGCCGGGTACCAGTCCTCTGCTCCCCAGGGAAACTGAACTCAGTTT	270 410
Db 290 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT	Db 350 GTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTT 409

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RESULT 22
US-10-216-167-15
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                                                                                              ; TYPE: DNA; ORGANISM: Homo Sapien US-10-216-167-15
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PRIOR FILLING DATE: 2002-04-09
PRIOR PELICATION NUMBER: 60/059113
PRIOR FILLING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
                               Query Match
Best Local Similarity
Matches 315; Conserv
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                                                                                                                                          SEQ ID NO 15
LENGTH: 1524
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APPLICANT:
APPLICANT:
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                                                                                                                                                                       Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 246
                                                                                                                                                                                                         PRIOR FILING DATE: 1998-03-27
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30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 89
                                                                                                                                                                                                                     FILING DATE: 1998-0.
APPLICATION NUMBER:
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APPLICATION NUMBER: 60/079294
FILING DATE: 1998-03-25
APPLICATION NUMBER: 60/079656
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FILING DATE: 1997-12-17
APPLICATION NUMBER: 60/078910
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Smith, Victoria
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Grimaldi, J. Christopher
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Gerritsen, Mary
Goddard, Audrey
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                                 Conservative
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                                              85.4%;
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                               Score 315; DB; Pred. No. 2e-
0; Mismatches
                                 0;
                               ; DB 42;
. 2e-159;
ches 0;
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                                                             Length 1524;
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; TYPE: DNA
; ORGANISM: Hor
US-10-216-168-15
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US-10-216-168-15
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PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
                                                                                                               NUMBER OF SEQ ID SEQ ID NO 15
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CURRENT FILING DATE: 2002-08-09
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                                                                                             LENGTH: 1524
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OR FILING DATE: 1998-03-20
OR APPLICATION NUMBER: 60/079294
OR FILING DATE: 1998-03-25
OR APPLICATION NUMBER: 60/079656
OR FILING DATE: 1998-03-26
OR FILING DATE: 1998-03-26
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                                                       Homo
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Smith, Victoria
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Grimaldi, J. Christopher
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Gerritsen, Mary
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85.48;
Score 315;
DB 42;
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Length 1524;
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US-10-218-612-15
                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
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GENERAL I
                   SEQ ID NO 15
LENGTH: 15
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                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/218,612
CURRENT FILING DATE: 2002-08-12
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR PPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR PPLICATION NUMBER: 60/062287
PRIOR PPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: WOOG, WIlliam I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530P1C21
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APPLICANT: Desno
                                                         Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 246
                                                                                           PRIOR FILING DATE: 1998-03-27
                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/063549 PRIOR FILING DATE: 1997-10-28
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 TYPE: DNA
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                                                                                                          FILING DATE: 1998-03-26
APPLICATION NUMBER: 60/079728
                                                                                                                                                      APPLICATION NUMBER: 60/079656
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTGCCTCGGCCCTCA 344
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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); Mismatches 0;
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                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/218,631
CURRENT FILING DATE: 2002-08-12
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/05287
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
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Best Local :
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                                                                                       APPLICATION NUMBER: 60/078910 FILING DATE: 1998-03-20 APPLICATION NUMBER: 60/079294 FILING DATE: 1998-03-25
                                                                                                                                                                                                           APPLICATION NUMBER: 60/064103 FILING DATE: 1997-10-31
                                   FILING DATE: 1998-03-26
APPLICATION NUMBER: 60/079728
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                                                                       APPLICATION NUMBER: 60/079656
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Prior Application data removed -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stephan, Jean-Philippe Watanabe, Colin L. Wood, William I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gurney, Austin L. Smith, Victoria
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Pred. No.
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See File Wrapper or PALM
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; ORGANISM: Homo
US-10-218-631-15
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US-10-218-765-15
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                                                   CURRENT APPLICATION NUMBER: US/10/218,765
CURRENT FILING DATE: 2002-08-12
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/06287
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
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PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
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SEQ ID NO 15
LENGTH: 1524
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Best Local Similarity
Matches 315; Conserv
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530P1C19
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                   APPLICATION NUMBER: 60/079294 FILING DATE: 1998-03-25
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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Gerritsen, Mary
Goddard, Audrey
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OR APPLICATION NUMBER: 60/089532
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APPLICATION NUMBER: 60/095916
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APPLICATION NUMBER: 60/096146
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APPLICATION NUMBER: 60/081955
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APPLICATION NUMBER: 60/096791
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APPLICATION NUMBER: 60/097986
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FILING DATE: 1998-09-10
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APPLICATION NUMBER: 60/134287
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APPLICATION NUMBER: 60/127887
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APPLICATION NUMBER: 60/101922
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APPLICATION NUMBER: 60/106178
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Best Local Similarity
CURRENT APPLICATION NUMBER: US/10/218,784 CURRENT FILING DATE: 2002-08-12 PRIOR APPLICATION NUMBER: 10/119,480 PRIOR FILING DATE: 2002-04-09 PRIOR APPLICATION NUMBER: 60/059113 PRIOR FILING DATE: 1997-09-17
                                                                                                   APPLICANT: Stephan, Jean-Philippe F.
APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530P1C18
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OR FILING DATE: 1999-08-17
OR APPLICATION NUMBER: 60/151733
OR FILING DATE: 1999-08-31
OR APPLICATION NUMBER: 60/164418
OR FILING DATE: 1999-11-09
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APPLICATION NUMBER: 60/146963
FILING DATE: 1999-08-03
APPLICATION NUMBER: 60/149320
FILING DATE: 1999-08-17
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FILING DATE: 1999-12-07
APPLICATION NUMBER: 60/169835
FILING DATE: 1999-12-07
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Watanabe, Colin L.
Wood, William I.
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Smith, Victoria
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Grimaldi, J. Christopher
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Gerritsen, Mary
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Pred. No. 2e-159;
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Sequence 15, Application US/10218849
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Desnoyers, Luc
APPLICANT: Gerritsen, Mary
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Grimaldi, J. Christopher
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DR FILING DATE: 1999-08-03
DR APPLICATION NUMBER: 60/149320
DR FILING DATE: 1999-08-17
DR APPLICATION NUMBER: 60/149638
DR FILING DATE: 1999-08-17
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FILING DATE: 1999-12-07
APPLICATION NUMBER: 60/169495
FILING DATE: 1999-12-07
APPLICATION NUMBER: 60/169835
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FILING DATE: 1999-11-16
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FILING DATE: 1999-08-31
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FILING DATE: 1999-07-28
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APPLICATION NUMBER: 60/141037
FILING DATE: 1999-06-23
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        Desnoyers, Luc
Gerritsen, Mary
Goddard, Audrey
Godowski, Paul J.
Grimaldi, J. Christopher
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Gurney, Austin L
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; ORGANISM: Homo Sapien US-10-218-930-15
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                              Prior Application removed -
NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
TYPE: DNA
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APPLICANT: Desnoyers, Lu
APPLICANT: Geritsen, Ma
APPLICANT: Goddard, Audr
APPLICANT: Godowski, Pau
APPLICANT: Grimaldi, J.
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APPLICANT:
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LENGTH: 1524
TYPE: DNA
ORGANISM: Homo Sapien
                                                                                                                   CURRENT APPLICATION NUMBER: US/10/218,930 CURRENT FILING DATE: 2002-08-12
                                                                                                                                                     TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3530P1C13
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CURRENT FILING DATE: 2002-08-12
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NUMBER OF SEQ ID NOS: 246
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                                                                                                                                                                                                             Stephan, Jean-Philippe F. Watanabe, Colin L. Wood, William I.
                                                                                                                                                                                                                                                                                                                                          Desnoyers, Luc
Gerritsen, Mary
Goddard, Audrey
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Grimaldi, J. Christopher
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                                                                                                    See File Wrapper or
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PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/064103
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Best Local Similarity
Matches 315; Conserv
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CURRENT FILING DATE: 2002-08-12
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                                        FILING DATE: 1998-03-26
APPLICATION NUMBER: 60/079728
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                                                                                         APPLICATION NUMBER: 60/
                                                                                                                   APPLICATION NUMBER: 60/079294
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           APPLICATION NUMBER: 60/081819 FILING DATE: 1998-04-15
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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Goddard, Audrey
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                                                                                          60/079656
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DR FILING DATE: 1998-06-25
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DR APPLICATION NUMBER: 60/095302
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APPLICATION NUMBER: 60/085579
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FILING DATE: 1998-06-18
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DR APPLICATION NUMBER: 60/130232
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APPLICATION NUMBER: 60/144758
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APPLICATION NUMBER: 60/141037
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APPLICATION NUMBER: 60/134287
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APPLICATION NUMBER: 60/127887
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FILING DATE: 1999-02-10
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FILING DATE: 1998-12-23
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APPLICATION NUMBER: 60/108849
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FILE REFERENCE: P3530PIC12

CURRENT APPLICATION NUMBER: US/10/219,003

CURRENT FILING DATE: 2002-08-12

PRIOR APPLICATION NUMBER: 10/119,480

PRIOR FILING DATE: 2002-04-09

PRIOR FILING DATE: 2002-04-09

PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/05287

PRIOR APPLICATION NUMBER: 60/06287

PRIOR FILING DATE: 1997-10-17

PRIOR APPLICATION NUMBER: 60/063549

PRIOR APPLICATION NUMBER: 60/063549

PRIOR APPLICATION NUMBER: 60/064103

PRIOR FILING DATE: 1997-10-28

PRIOR APPLICATION NUMBER: 60/064103

PRIOR FILING DATE: 1997-10-31
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US-10-219-003-15
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GENERAL I
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PRIOR APPLICATION NUMBER: 60/151733
PRIOR APPLICATION NUMBER: 60/164418
PRIOR APPLICATION NUMBER: 60/164418
PRIOR FILING DATE: 1999-11-09
PRIOR FILING DATE: 1999-11-16
PRIOR FILING DATE: 1999-11-16
PRIOR APPLICATION NUMBER: 60/16361
PRIOR APPLICATION NUMBER: 60/169445
PRIOR FILING DATE: 1999-12-07
PRIOR APPLICATION NUMBER: 60/169495
PRIOR APPLICATION NUMBER: 60/169495
PRIOR APPLICATION NUMBER: 60/169495
PRIOR APPLICATION NUMBER: 60/169495
PRIOR APPLICATION NUMBER: 60/169495
PRIOR APPLICATION NUMBER: 60/169495
PRIOR FILING DATE: 1999-12-07
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Baker, Kevin P.
APPLICANT: Desnoyers, Luc
APPLICANT: Gernitsen, Mary
APPLICANT: Goddard, Audrey
APPLICANT: Goddowski, Paul I.
APPLICANT: Grimaldi, J. Chi
APPLICANT: Gurney, Austin I
                                                                                                                                                                                                                                                                               APPLICANT: Stephan, Jean-Philippe F.
APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
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Godowski, Paul J.
Gorimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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Gerritsen, Mary
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1998-0 1998-0 1998-0 NUMBER: 1998-0 NUMBER: 1998-0 NUMBER: 1998-0 NUMBER: 1998-0 NUMBER:	PRIOR APPLICATION NUMBER: 60/095302 PRIOR APPLICATION NUMBER: 60/095318 PRIOR APPLICATION NUMBER: 60/095318 PRIOR APPLICATION NUMBER: 60/095916 PRIOR APPLICATION NUMBER: 60/095916 PRIOR APPLICATION NUMBER: 60/096146 PRIOR FILING DATE: 1998-08-11 PRIOR APPLICATION NUMBER: 60/096791 PRIOR APPLICATION NUMBER: 60/096791 PRIOR APPLICATION NUMBER: 60/097986 PRIOR FILING DATE: 1998-08-17 PRIOR FILING DATE: 1998-08-26 PRIOR FILING DATE: 1998-08-26 PRIOR APPLICATION NUMBER: 60/09594 PRIOR APPLICATION NUMBER: 60/09596 PRIOR APPLICATION NUMBER: 60/09596 PRIOR APPLICATION NUMBER: 60/09598 PRIOR APPLICATION NUMBER: 60/09598 PRIOR APPLICATION NUMBER: 60/09598 PRIOR APPLICATION NUMBER: 60/09598 PRIOR APPLICATION NUMBER: 60/09598 PRIOR APPLICATION NUMBER: 60/09598 PRIOR APPLICATION NUMBER: 60/09598	NUMBER: 60/08639 11998-05-22 11998-06-17 11998-06-17 NUMBER: 60/08953 11998-06-17 NUMBER: 60/08953 11998-06-18 NUMBER: 60/09047 NUMBER: 60/09059 11998-06-25 11998-06-25 11998-06-25 NUMBER: 60/09059 11998-06-25 NUMBER: 60/09059	PRIOR APPLICATION NUMBER: 60/069873 PRIOR APPLICATION NUMBER: 60/078910 PRIOR APPLICATION NUMBER: 60/078910 PRIOR APPLICATION NUMBER: 60/079294 PRIOR APPLICATION NUMBER: 60/079294 PRIOR APPLICATION NUMBER: 60/079656 PRIOR APPLICATION NUMBER: 60/079728 PRIOR EILING DATE: 1998-03-26 PRIOR APPLICATION NUMBER: 60/079728 PRIOR APPLICATION NUMBER: 60/081819 PRIOR APPLICATION NUMBER: 60/081819 PRIOR FILING DATE: 1998-04-15 PRIOR APPLICATION NUMBER: 60/081955 PRIOR APPLICATION NUMBER: 60/082804 PRIOR APPLICATION NUMBER: 60/082804 PRIOR APPLICATION NUMBER: 60/082804 PRIOR APPLICATION NUMBER: 60/082804 PRIOR APPLICATION NUMBER: 60/082804 PRIOR APPLICATION NUMBER: 60/082804 PRIOR APPLICATION NUMBER: 60/08323 PRIOR APPLICATION NUMBER: 60/085579 PRIOR APPLICATION NUMBER: 60/085579
PRIOR APPLICATION NUMBER: 60/131022 PRIOR FILING DATE: 1999-04-26 PRIOR APPLICATION NUMBER: 60/131270 PRIOR FILING DATE: 1999-04-27 PRIOR FILING DATE: 1999-04-27 PRIOR FILING DATE: 1999-04-27 PRIOR FILING DATE: 1999-04-27 PRIOR APPLICATION NUMBER: 60/131445 PRIOR FILING DATE: 1999-04-28 PRIOR APPLICATION NUMBER: 60/134287 PRIOR FILING DATE: 1999-05-14 PRIOR APPLICATION NUMBER: 60/140650 PRIOR APPLICATION NUMBER: 60/140723 PRIOR APPLICATION NUMBER: 60/140723 PRIOR APPLICATION NUMBER: 60/140723 PRIOR APPLICATION NUMBER: 60/140723	PRIOR FILING DATE: 1999-01-12 PRIOR APPLICATION NUMBER: 60/115565 PRIOR APPLICATION NUMBER: 60/115733 PRIOR APPLICATION NUMBER: 60/115733 PRIOR FILING DATE: 1999-01-12 PRIOR APPLICATION NUMBER: 60/119549 PRIOR FILING DATE: 1999-02-10 PRIOR APPLICATION NUMBER: 60/123618 PRIOR APPLICATION NUMBER: 60/12559 PRIOR APPLICATION NUMBER: 60/12575 PRIOR APPLICATION NUMBER: 60/125775 PRIOR APPLICATION NUMBER: 60/125775 PRIOR APPLICATION NUMBER: 60/126773 PRIOR APPLICATION NUMBER: 60/126773 PRIOR APPLICATION NUMBER: 60/127887 PRIOR APPLICATION NUMBER: 60/127887 PRIOR APPLICATION NUMBER: 60/127887 PRIOR APPLICATION NUMBER: 60/127887 PRIOR APPLICATION NUMBER: 60/127887 PRIOR APPLICATION NUMBER: 60/127887 PRIOR APPLICATION NUMBER: 60/130232 PRIOR APPLICATION NUMBER: 60/130232	PRIOR APPLICATION NUMBER: 60/106464 PRIOR FILING DATE: 1998-10-30 PRIOR FILING DATE: 1998-10-30 PRIOR FILING DATE: 1998-11-03 PRIOR FILING DATE: 1998-11-07 PRIOR FILING DATE: 1998-11-17 PRIOR FILING DATE: 1998-11-17 PRIOR APPLICATION NUMBER: 60/108801 PRIOR APPLICATION NUMBER: 60/108849 PRIOR FILING DATE: 1998-11-18 PRIOR PILING DATE: 1998-11-18 PRIOR PILING DATE: 1998-12-15 PRIOR APPLICATION NUMBER: 60/112422 PRIOR FILING DATE: 1998-12-23 PRIOR FILING DATE: 1998-12-23 PRIOR FILING DATE: 1998-12-23 PRIOR FILING DATE: 1998-12-23 PRIOR PILING DATE: 1998-12-33 PRIOR APPLICATION NUMBER: 60/113621 PRIOR APPLICATION NUMBER: 60/113621 PRIOR APPLICATION NUMBER: 60/113621 PRIOR APPLICATION NUMBER: 60/113621 PRIOR APPLICATION NUMBER: 60/113621	PRIOR FILING DATE: 1998-09-16 PRIOR APPLICATION NUMBER: 60/100848 PRIOR APPLICATION NUMBER: 60/100919 PRIOR APPLICATION NUMBER: 60/100919 PRIOR FILING DATE: 1998-09-17 PRIOR APPLICATION NUMBER: 60/101477 PRIOR APPLICATION NUMBER: 60/101738 PRIOR APPLICATION NUMBER: 60/101738 PRIOR FILING DATE: 1998-09-24 PRIOR FILING DATE: 1998-09-24 PRIOR FILING DATE: 1998-09-24 PRIOR APPLICATION NUMBER: 60/101746 PRIOR APPLICATION NUMBER: 60/101916 PRIOR FILING DATE: 1998-09-25 PRIOR APPLICATION NUMBER: 60/101922 PRIOR FILING DATE: 1998-09-24 PRIOR FILING DATE: 1998-09-24 PRIOR FILING DATE: 1998-09-24 PRIOR APPLICATION NUMBER: 60/101922 PRIOR APPLICATION NUMBER: 60/106178 PRIOR APPLICATION NUMBER: 60/106178 PRIOR APPLICATION NUMBER: 60/106178 PRIOR APPLICATION NUMBER: 60/106248 PRIOR FILING DATE: 1998-10-29

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RESULT 32
US-10-219-010-15
Sequence 15, Applicati
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL DESNOYER
APPLICANT: Baker, Key
APPLICANT: DESNOYER
APPLICANT: Goddard,
APPLICANT: Goddwski
APPLICANT: Godowski
APPLICANT: Godowski
APPLICANT: Gurney,
APPLICANT: Smith, V.
APPLICANT: Stephan,
APPLICANT: Watanabe,
APPLICANT: Watanabe,
APPLICANT: Wood, Wil
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OR FILING DATE: 1999-06-23
OR APPLICATION NUMBER: 60/144758
OR FILING DATE: 1999-07-20
OR APPLICATION NUMBER: 60/145698
OR FILING DATE: 1999-07-28
OR FILING DATE: 1999-07-28
OR APPLICATION NUMBER: 60/146963
OR FILING DATE: 1999-08-03
OR APPLICATION NUMBER: 60/149320
OR FILING DATE: 1999-08-17
OR APPLICATION NUMBER: 60/15173
OR FILING DATE: 1999-08-17
OR APPLICATION NUMBER: 60/15173
OR FILING DATE: 1999-08-17
OR APPLICATION NUMBER: 60/164418
OR FILING DATE: 1999-11-09
OR APPLICATION NUMBER: 60/166361
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   INVENTION:
                 INVENTION:
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                                                       Godowski, Paul J.
Grinaldi, J. Christopher
Gurney, Austin I.
Smith, Victoria
Stephan, Jean-Philippe F.
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Gerritsen, Mary
Goddard, Audrey
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                             Watanabe, Colin L. Wood, William I.
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ilarity 100.0%;
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CURRENT APPLICATION NUMBER:
CURRENT FILING DATE: 2002.
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IOR APPLICATION NUMBER: 60/141037
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APPLICATION NUMBER: 60/134287
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APPLICATION NUMBER: 60/
FILING DATE: 1999-04-27
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FILING DATE: 1999-07-26
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FILING DATE: 1999-07-20
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RESULT 33
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; Sequence 15, Application US/10219060
; GENERAL INFORMATION:

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PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
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CURRENT FILING DATE: 2002-08-12
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              OR APPLICATION NUMBER: 60/090691
OR APPLICATION NUMBER: 60/090691
OR APPLICATION NUMBER: 60/090695
OR APPLICATION NUMBER: 60/09095
OR FILING DATE: 1998-06-25
OR APPLICATION NUMBER: 60/091982
OR FILING DATE: 1998-07-07
OR APPLICATION NUMBER: 60/095302
OR FILING DATE: 1998-08-04
OR APPLICATION NUMBER: 60/095318
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OR FILING DATE: 1998-06-17
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DR FILING DATE: 1998-05-13
DR APPLICATION NUMBER: 60/085579
DR FILING DATE: 1998-05-15
DR APPLICATION NUMBER: 60/086392
DR FILING DATE: 1998-05-22
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OR FILING DATE: 1998-03-26
OR APPLICATION NUMBER: 60/079728
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OR FILING DATE: 1998-04-15
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APPLICATION NUMBER: 60/090557
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APPLICATION NUMBER: 60/082804
FILING DATE: 1998-04-22
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FILING DATE: 1998-05-06
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Gerritsen, Mary
Goddard, Audrey
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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OR APPLICATION NUMBER: 60//13605
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OR APPLICATION NUMBER: 60//13621
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OR APPLICATION NUMBER: 60//11558
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OR FILING DATE: 1999-01-12
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OR APPLICATION NUMBER: 60/100919
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OR FILING DATE: 1998-09-16 APPLICATION NUMBER: 60/112422 FILING DATE: 198-12-15 APPLICATION NUMBER: 60/113296 FILING DATE: 1998-12-22 FILING DATE: 1998-10-29 APPLICATION NUMBER: 60/1 FILING DATE: 1998-10-30 APPLICATION NUMBER: 60/106178 FILING DATE: 1998-10-28 APPLICATION NUMBER: 60/106248 FILING DATE: 1998-11-APPLICATION NUMBER: FILING DATE: 1998-08-17 APPLICATION NUMBER: 60/ FILING DATE: 1998-08-26 APPLICATION NUMBER: 60/098544 FILING DATE: 1998-08-31 APPLICATION NUMBER: 60/ 1998-09-16 1998-09-16 ""MHER: 60/100848 60/108849 60/106464 60/097986 60/096791

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FILING DATE: 1999-04-27
APPLICATION NUMBER: 60/131445
FILING DATE: 1999-04-28
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FILING DATE: 1999-04-26
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FILING DATE: 1999-12-07
APPLICATION NUMBER: 60/169835
FILING DATE: 1999-12-07
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APPLICATION NITTON
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APPLICATION NUMBER: 60/140723
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FILING DATE: 1999-11-16
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FILING DATE: 1999-12-07
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CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 229
                                                                                                                     AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
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                                                              AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT
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CURRENT FILING DATE: 2002-08-12
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/06287
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/063549
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SEQ ID NO 15
LENGTH: 1524
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Best Local
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PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
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PRIOR APPLICATION NUMBER: 60/079728
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PRIOR APPLICATION NUMBER: 60/079294
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                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 1998-03-27
                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Homo Sapien
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                  AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
                                                                                                                                       Similarity
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Grimaldi, J. Christopher
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Gerritsen, Mary
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                                                                                                                                                                                                         85.4%; Score 315; DE 100.0%; Pred. No. 2e-tive 0; Mismatches
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; ORGANISM: Homo Sapien
US-10-219-062-15
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
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PRIOR APPLICATION NUMBER: 60/062287
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                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/079728 PRIOR FILING DATE: 1998-03-27
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CURRENT FILING DATE: 2002-08-12
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230 GCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 289
                                                                    170 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 229
                  90 GCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAGAAAG 149
                                                                                    30 CAGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACT 89
                                                                                                                                                                                                                                                                                                                                                                                                                                    OR APPLICATION NUMBER: 60/063549
OR ETLING DATE: 1997-10-28
OR APPLICATION NUMBER: 60/064103
OR FILING DATE: 1997-10-31
OR APPLICATION NUMBER: 60/069873
OR ETLING DATE: 1997-12-17
OR APPLICATION NUMBER: 60/078910
OR ETLING DATE: 1998-03-20
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FILING DATE: 1998-03-25
APPLICATION NUMBER: 60/079656
FILING DATE: 1998-03-26
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                                                                                                                                                                Similarity
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Gerritsen, Mary
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                                                                                                                                                Conservative
                                                                                                                                         100.0%; **
                                                                                                                                                            85.4%; Score 315; 100.0%; Pred. No.
                                                                                                                                            Mismatches
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LENGTH: 1524
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Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 246
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR ETILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
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PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
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Local Similarity
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Grimaldi, J. Christopher
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Gerritsen, Mary
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                                                                                Conservative
                                                                        85.4%; Score 315; DB 42;
100.0%; Pred. No. 2e-159;
... Mismatches 0;
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                                                                                                            Length 1524;
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PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR TILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/06287
PRIOR TILING DATE: 1997-10-17
PRIOR PELLING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/06349
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/069873
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NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
       Query Match
Best Local Similarity
Matches 315; Conserv
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CURRENT FILING DATE: 2002-08-13
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ORGANISM: HOMO
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Grimaldi, J. Christopher
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       Conservative
                           85.4%; Score 315; DB 42; 100.0%; Pred. No. 2e-159;
     0;
  Mismatches
                                                  DB 42;
                                                                                                                                                                                                                                         See File Wrapper or PALM
     0;
                                             Length 1524;
  Indels
  0,
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US-10-219-065-15; Sequence 15; Apr
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PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2022-04-09
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/06287
PRIOR APPLICATION NUMBER: 60/063849
PRIOR FILING DATE: 1997-10-28
PRIOR PAPPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-20
PRIOR FILING DATE: 1998-03-25
PRIOR FILING DATE: 1998-03-25
PRIOR FILING DATE: 1998-03-25
PRIOR FILING DATE: 1998-03-25
PRIOR FILING DATE: 1998-03-26
                                                              Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15, Application US/10219065 GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER: US/10/219,065
CURRENT FILING DATE: 2002-08-13
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                                                                                                                          PRIOR APPLICATION NUMBER: 60/079728 PRIOR FILING DATE: 1998-03-27
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TYPE: DNA
ORGANISM: Homo Sapien
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                                               LENGTH: 1524
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                                                                                                                                                               APPLICATION NUMBER: 60/079656
FILING DATE: 1998-03-26
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Gerritsen, Mary
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                                                                                                   See File Wrapper or PALM
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                                                                            CURRENT APPLICATION NUMBER: US/10/219,066
CURRENT FILING DATE: 2002-08-13
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064873
PRIOR APPLICATION NUMBER: 60/079910
PRIOR APPLICATION NUMBER: 60/079910
PRIOR APPLICATION NUMBER: 60/079910
PRIOR APPLICATION NUMBER: 60/079910
PRIOR APPLICATION NUMBER: 60/079910
PRIOR APPLICATION NUMBER: 60/079994
PRIOR FILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/079656
PRIOR APPLICATION NUMBER: 60/079656
PRIOR APPLICATION NUMBER: 60/079656
PRIOR APPLICATION NUMBER: 60/079656
PRIOR FILING DATE: 1998-03-26
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APPLICANT:
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                                             PRIOR APPLICATION NUMBER: 60/079656
PRIOR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/079728
PRIOR FILING DATE: 1998-03-27
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                Remaining Prior Application NUMBER OF SEQ ID NOS: 246
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ID NO 1
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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Gerritsen, Mary
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                                    removed
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                                    File
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US-10-219-067-15
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US-10-219-066-15
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                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR PEPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-28
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                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: P3530P1C51
CURRENT APPLICATION NUMBER: US/10/219,067
CURRENT FILING DATE: 2002-08-14
                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND
TITLE OF INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                          PRIOR PRIOR
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                                                                                                                       PRIOR
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nes 315; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90
                                                                                                   APPLICATION NUMBER: 60/069873
FILING DATE: 1997-12-17
APPLICATION NUMBER: 60/078910
                                                                                                                                                          APPLICATION NUMBER: 60/063549 FILING DATE: 1997-10-28 APPLICATION NUMBER: 60/064103 FILING DATE: 1997-10-31
                                  FILING DATE: 1998-0: APPLICATION NUMBER:
                                                                   APPLICATION NUMBER: 60/079294
                                                                                     FILING DATE:
                   FILING DATE:
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Smith, Victoria
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Grimaldi, J. Christopher
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Gerritsen, Mary
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   NUMBER:
                     1998-03-26
                                                      1998-03-25
                                                                                       1998-03-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Austin L.
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100.0%;
                                      60/079656
   60/079728
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; Remaining Prior Application data removed -
; NUMBER OF SEQ ID NOS: 246
; SEQ ID NO 15
; LENGTH: 1524
; TYPE: DNA
; ORGANISM: Homo Sapien
US-10-219-067-15
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                                   FILE REFERENCE: P3530FLC31
CURRENT APPLICATION NUMBER: US/10/219,068
CURRENT EILING DATE: 2002-08-13
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/06287
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/06403
PRIOR FILING DATE: 1997-10-31
PRIOR PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
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APPLICATION NUMBER: 60/079294 FILING DATE: 1998-03-25
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Godowski, Paul J
Grimaldi, J. Christopher
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Gerritsen, Mary
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PRIOR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/079728
PRIOR FILING DATE: 1998-03-27
Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 15, Applica GENERAL INFORMATION:
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Best Local Similarity
Matches 315; Conserv
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APPLICANT:
                                                                                                                            CURRENT APPLICATION NUMBER: US/10/219, CURRENT FILING DATE: 2002-08-13 PRIOR APPLICATION NUMBER: 10/119,480 PRIOR EILING DATE: 2002-04-09 PRIOR APPLICATION NUMBER: 60/059113 PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                      TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3530P1C40
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
APPLICANT:
                                                    PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-28
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                                                  APPLICATION NUMBER: 60/063549 FILING DATE: 1997-10-28
               APPLICATION NUMBER: 60/064103 FILING DATE: 1997-10-31
APPLICATION NUMBER: 60/069873
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Watanabe, Colin L.
Wood, William I.
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Gerritsen, Mary
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530P1C42
CURRENT APPLICATION UMBER: US/10/219,070
CURRENT FILING DATE: 2002-08-13
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/05287
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/063549
PRIOR FILING DATE: 1997-10-17
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APPLICANT: Godowski, Paul J.
APPLICANT: Godowski, Paul J. Christopher
APPLICANT: Gurney, Austin L.
APPLICANT: Smith, Victoria
APPLICANT: Smith, Victoria
APPLICANT: Watanabe, Colin L.
APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, Milliam I.
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PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/079656
PRIOR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/079728
PRIOR FILING DATE: 1998-03-27
PRIOR FILING DATE: 1998-03-27
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Best Local S
Matches 315
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APPLICANT: Desnoye
APPLICANT: Gerrits
APPLICANT: Goddard
APPLICANT: Godowsk
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SEQ ID NO 15
LENGTH: 1524
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les 315; Conservative 0; Mismatches
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Gerritsen, Mary
                                                                                                                                                                                                                                                                                                                                                                                                                                     Kevin
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2e-159;
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APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Smith, Victoria
APPLICANT: Stephan, Jean-Philippe F.
APPLICANT: Watanabe, Colin L.
APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCL
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3530P1C46
CURRENT APPLICATION NUMBER: US/10/219,071
CURRENT FILING DATE: 2002-08-13
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 2002-04-09
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                                                                                                                                                                                                                                                           APPLICANT: Baker, Kevin P.
APPLICANT: Desnoyers, Luc
APPLICANT: Gerritsen, Mary
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul
APPLICANT: Grimaldi, J. Chi
APPLICANT: Grimaldi, J. Chi
APPLICANT: Grimaldi, J. Chi
APPLICANT: Grimaldi, J. Chi
APPLICANT: Smith, Victoria
APPLICANT: Stephan, Jean-Pi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15, Application US/10219071 GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
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PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
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PRIOR APPLICATION NUMBER: 60/059113 PRIOR FILING DATE: 1997-09-17
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Gerritsen, Mary
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100.0%; Pred. No. 2e:
tive 0; Mismatches
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US-10-219-072-15
Sequence 15, Application US/10219072
SEQUENCE 15, Application US/10219072
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Besnoyers, Luc
APPLICANT: Gerritsen, Mary
APPLICANT: Gerritsen, Mary
APPLICANT: Goddard, Audrey
APPLICANT: Goddowski, Paul J.
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Gurney, Austin L.
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PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/079656
PRIOR APPLICATION NUMBER: 60/079728
PRIOR APPLICATION NUMBER: 60/079728
PRIOR PILING DATE: 1998-03-27
PRIOR FILING DATE: 1998-03-27
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NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
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PRIOR
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3530P1C45 CURRENT APPLICATION NUMBER: US/10/219,072 CURRENT FILING DATE: 2002-08-13
                                                                                                  APPLICANT:
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                                                                                                                     APPLICANT:
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R FILING DATE: 1997-10-28

R FILING DATE: 1997-10-31

DR APPLICATION NUMBER: 60/069873

DR APPLICATION NUMBER: 60/069873

DR FILING DATE: 1997-12-17
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                                                                                                Goddard, Audrey
Goddwski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
Stephan, Jean-Philippe F.
Watanabe, Colin L.
Wood, William I.
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| MBRANE<br>SAME<br>- See                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            |
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| POLYF                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         | 918<br>94<br>94<br>94<br>94                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |
| Wrag                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            |
| POLYPEPTIDES                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  | 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      |
| Or                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            | sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence sequence                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        |
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| NUCLEIC                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | 48790<br>48791<br>48792<br>48792<br>94484<br>94485<br>94488<br>94488<br>94488<br>16179<br>147785<br>13316<br>18596<br>18527<br>28218<br>25794                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |

Query Match

85.4%;

Score 315;

DB 6;

Length 1524;

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US-10-092-411A-2083/C

US-10-092-411A-2083/C

Sequence 2083, Application US/10092411A

GENERAL INFORMATION:
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; ORGANISM: Staphylococcus epidermidis
US-10-092-411A-1425
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CURRENT FILING DATE: 2002-03-07
PRIOR APPLICATION NUMBER: US 09/134,001
PRIOR FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5676
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Matches 315; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1425, Application US/10092411A GENERAL INFORMATION:
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LENGTH: 1587
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Lynn Doucette-Stamm et al TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: 032796-101
PRIOR APPLICATION NUMBER: US 60/064,964
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             567 AATTCCAGCTGAACAACG 584
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.9%; Score 18; DB 6; Length 1587; 100.0%; Pred. No. 0.74; tive 0; Mismatches 0; Indels
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PRIOR FILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/064,964
PRIOR FILING DATE: 1997-11-08
PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5676
SEQ ID NO 263
LENGTH: 393
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LENGTH: 1596
TYPE: DNA
ORGANISM: Staphylococcus epidermidis
US-10-092-411A-2083
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                                                                   ; TYPE: DNA; ORGANISM: Homo sapiens US-09-513-999C-36268
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; GENERAL INFORMATION:
; APPLICANT: Dumas Milne Edwards, J.B.
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Matches
                                                                                                                                                       NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 36268
LENGTH: 111
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PRIOR APPLICATION NUMBER: US 60/055,779
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 5676
      Query Match
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CURRENT APPLICATION NUMBER: US/10/092,411A
CURRENT FILING DATE: 2002-03-07
                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24
                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/122,487 PRIOR FILING DATE: 1999-02-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1308 TGCATCAGCTGCTGCAAC 1291
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18; Conserv
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            4.38;
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            Score 16;
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Best Local Similarity
Matches 16; Conser
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LENGTH: 397
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APPLICANT: Dumas Milne Edwards, J.B.
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SOFTWARE: Patent.pm
SEQ ID NO 28601
LENGTH: 433
TYPE: DNA
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TITLE OF INVENTION: Novel Murine Polynucleotide Tags and
TITLE OF INVENTION: Mutant Cells and Mutant Animals Observe Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Processing Control of the Pro
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PRIOR FILING DATE: 1999-07-15
NUMBER OF SEQ ID NOS: 2908
SOFTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Zambrowicz, Bri
APPLICANT: Sands, Arthur T
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CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Duclert, A. APPLICANT: Giordano, J.Y. TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature LOCATION: 275
                                                                                                                                                              OTHER INFORMATION: r-a or 9
                                                                                                                                                                                                                              NAME/KEY: misc_feature
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                                      CN: 310
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TION: r-a or g
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US-09-513-999C-3777/c
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SOFTWARE: Patent.pm
SEQ ID NO 3777
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SEQ ID NO 24773
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TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Dumas Milne Edwards, J.B. APPLICANT: Duclert, A.
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NAME/KEY: sig_peptide
LOCATION: 73..123
OTHER INFORMATION: score 9.3
OTHER INFORMATION: seg LALSLLILVLAFG/IP
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ORGANISM: Homo sapiens
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LOCATION: 73..474
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                                                    NAME/KEY: misc_feature LOCATION: 182
                                                                                           ORGANISM: Homo sapiens FEATURE:
NAME/KEY: misc_feature
                  FEATURE:
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US-09-513-999C-14500/c

; Sequence 14500, Application US/09513999C

; GENERAL INFORMATION:
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; OTHER INFORMATION: k-g or t
US-09-513-999C-24773
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Best Local Similarity
                   APPLICANT: Dumas Milne Edwards, J.B. APPLICANT: Duclert, A.
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GENERAL INFORMATION:
      APPLICANT:
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PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
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LOCATION: 197
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LOCATION: 141..464
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LOCATION: 198
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Giordano, J.Y.
                Duclert, A.
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100.0%; Pred. No.
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US-10-240-425-140
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LENGTH: 579
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Best Local S
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                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/240,425
CURRENT FILING DATE: 2002-09-30
PRIOR APPLICATION NUMBER: PCT/US01/09847
PRIOR FILING DATE: 2001-03-28
PRIOR APPLICATION NUMBER: US 60/193,446
PRIOR FILING DATE: 2000-03-31
NUMBER OF 5EQ ID NOS: 1588
SOFTWARE: Patentin Ver. 2.1
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SEQ ID NO 14500
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TITLE OF INVENTION: Gene Expression Profiles in
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APPLICANT: Boland, Joseph
APPLICANT: Lord, Reginald
APPLICANT: Alvarez, Chris
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                                                                                                                                                                     TYPE: DNA
ORGANISM: Homo sapiens
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LOCATION: 325
OTHER INFORMATION: r=a or
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OTHER INFORMATION: y=c
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81 GCCAAGGCCCAAGAA 95
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                                                                15;
                                                                               Similarity
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Scherf, Uwe
                                                               Conservative
                                                                             100.0%;
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100.0%; Pred. No.
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; LCCATION: (1)..(686)
; OTHER INFORMATION: n - a or c or g
US-10-240-425-710
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PRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 1588
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 710
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TITLE OF INVENTION: Gene Expression Profiles in Esophageal Tissue
FILE REFERENCE: 44921-5026
CURRENT APPLICATION NUMBER: US/10/240,425
CURRENT FILING DATE: 2002-09-30
PRIOR APPLICATION NUMBER: PCT/US01/09847
PRIOR FILING DATE: 2001-03-28
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Scherf, Uwe
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Lord, Reginald V.
Alvarez, Chris
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                                                                                                                                                                                             ELLIOTT, Vicki S. FORSYTHE, Ian J. BECHA, Shanya D.
                                                                                                                                                                                                                                           TANG, Y. Tom
CHAWLA, Narinder K.
                                                                                                                                                                                                                                                                                                     THANGAVELU, Kavitha
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                                                          LEE, Ernestine A.
LEE, Soo Yeun
ELLIOTT, Vicki S.
                                                                                                                                                  GRIFFIN, Jennifer A.
ĽÜ,
              GORVAD, Ann E
KABLE, Amy E.
                                                                                                       BAUGHN, Mariah R.
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                                            RAMKUMAR, Jayalaxmi
                                                                                                                       ZEBARJADIAN,
                                                                                                                                     LAL, Preeti G.
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Dyung Aina M.
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                                                                                       Ernestine A.
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100.0%;
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Pred. No.
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US-10-264-237-1045
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SEQ ID NO 35
LENGTH: 940
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Best Local Similarity
Matches 15; Conserv
                                   Sequence 1045, Application US/10264237 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Ver. 3.1
SEQ ID NO 561
LENGTH: 1292
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                                                                                                                                                                                                                                               Query Match
Best Local
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PRIOR FILING DATE: 2001-10-19
PRIOR APPLICATION NUMBER: US 60/343,980
PRIOR FILING DATE: 2001-11-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 2001-11-02
PRIOR APPLICATION NUMBER: US 60/332,426
PRIOR FILING DATE: 2001-11-16
NUMBER OF SEO ID NOS: 64
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TITLE OF INVENTION: SECRETED PROTEINS
FILE REFERENCE: PF-1217 PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 2876
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: PCT/US02/32032 CURRENT FILING DATE: 2002-10-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Nuc
FILE REFERENCE: PAl31P1
APPLICANT: Birse et
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Birse et al.
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                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
FEATURE:
REATURE:
NAME/KEY: misc_feature
LOCATION: (534)..(534)
OTHER INFORMATION: n equals a,t,g, or c
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                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/10264237
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Nucleic Acids, Proteins, and Antibodies
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100.0%; P1
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L00.0%; Pred. No. 34;
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Pred. No.
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US-10-125-923A-555/c ; Sequence 555, Application US/10125923A ; GENERAL IMFORMATION:
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                                                                                                            RESULT 18
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GENERAL INFORMATION:
APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PA131p1
CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
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LENGTH: 1733
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LENGTH: 1542
                   APPLICANT: Baker, Kevin P. APPLICANT: Chen, Jian
   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
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PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 2876
SOFTWARE: Patentin Ver. 3.1
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SOFTWARE: PatentIn Ver. 3.
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CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature
LOCATION: (1459)..(1459)
OTHER INFORMATION: n equals a,t,g, or c
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ORGANISM: Homo sapiens
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LOCATION: (1471)...(1471)
OTHER INFORMATION: n equals a,t,g, or
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100.0%; Pred. No.
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Pred. No.
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; ORGANISM: Homo sapiens US-10-264-237-1252
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                                                                                                                        SOFTWARE: PatentIn Ver.
SEQ ID NO 1252
LENGTH: 1913
Query Match
Best Local Similarity
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GENERAL INFORMATION:
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LENGTH: 1773
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CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
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TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PA131P1
                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 2876
                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 60/205,515 PRIOR FILING DATE: 2000-05-19
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PRIOR APPLICATION NUMBER: 60/063120
PRIOR FILING DATE: 1997-10-24
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PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059266
PRIOR FILING DATE: 1997-09-18
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                                                                                                       TYPE: DNA
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4.18;
100.08;
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Pred. No.
  Score
Pred.
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  . No.
  DB 6;
35;
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                   Length 1913;
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FILE REFERENCE: 475412001300
CURRENT APPLICATION NUMBER: US/10/192,280
CURRENT FILING DATE: 2002-07-09
NUMBER OF SEQ ID MOS: 39
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
LENGTH: 25020
TYPE: DNA
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US-10-240-851-69/c
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PRIOR FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 09/543,771
PRIOR FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 09/229,319
PRIOR FILING DATE: 1999-01-13
PRIOR APPLICATION NUMBER: US 60/071,449
PRIOR FILING DATE: 1998-01-13
PRIOR APPLICATION NUMBER: US 60/105,511
PRIOR APPLICATION NUMBER: US 60/105,511
PRIOR APPLICATION NUMBER: US 60/105,511
PRIOR FILING DATE: 1998-10-23
RESULT 22
US-10-240-425-1461
                                                                                                                                                                                                                       ; ORGANISM: Streptococcus agalactiae 
US-10-192-280-1
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GENERAL INFORMATION:
APPLICANT: Dattagupta, Nanibhushan
APPLICANT: Shah, Ketan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DETECTING
TITLE OF INVENTION: STREPTOCOCCUS AGALACTIAE CAPSULAR POLYSACCHARIDE SYNTHESIS
TITLE OF INVENTION: GENES
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SEQ ID NO 69
LENGTH: 3513
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                                                                                                                                            Query Match 4.1%;
Best Local Similarity 100.0%;
Matches 15; Conservative
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/240,851
CURRENT FILING DATE: 2002-10-04
                                                                     10941 AAGAAGTGATGGAGC 10955
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Pred. No.
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APPLICANT: Scherf, Uwe
APPLICANT: Vockley, Joseph G.
TITLE OF INVENTION: Gene Expression Profiles in Esophageal Tissue
FILE REFERENCE: 44921-5026
CURRENT APPLICATION NUMBER: US/10/240,425
CURRENT FILING DATE: 2002-09-30
PRIOR APPLICATION NUMBER: PCT/US01/09847
PRIOR APPLICATION NUMBER: PCT/US01/09847
PRIOR FILING DATE: 2001-03-28
PRIOR FILING DATE: 2001-03-28
PRIOR FILING DATE: 2000-03-31
VMMBER OF SEQ ID NOS: 1588
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1121
LENGTH: 74822
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                                                                                                                                                 ; OTHER INFORMATION: Genbank Accession No. AL049798 US-10-240-425-1121
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PRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 1588
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 1461
LENGTH: 36534
TYPE: DNA
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Best Local S
Matches 15
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GENERAL INFORMATION:
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Best Local Similarity
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APPLICANT: Scherf, Uwe
APPLICANT: Vockley, Joseph G.

TITLE OF INVENTION: Gene Expression Profiles in Esophageal Tissue
FILE REFERENCE: 44921-5026
CURRENT APPLICATION NUMBER: US/10/240,425
CURRENT FILING DATE: 2002-09-30
PRIOR APPLICATION NUMBER: PCT/US01/09847
PRIOR FILING DATE: 2001-03-28
PRIOR FILING DATE: 2001-03-28
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APPLICANT:
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APPLICANT: Boland, Joseph I
APPLICANT: Lord, Reginald I
APPLICANT: Alvarez, Chris
APPLICANT: Wetzel, Jon C.
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70056 ATGTGTCAGAAAGAA 70070
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                   137 ATGTGTCAGAAAGAA 151
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                                                                       Local Similarity 100 nes 15; Conservative
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Lord, Reginald V.
Alvarez, Chris
Wetzel, Jon C.
Scherf, Uwe
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Lord, Reginald V.
Alvarez, Chris
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                                                                                         4.1%; Score 15;
100.0%; Pred. No.
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Pred. No.
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US-60-417-190-48787/c
Sequence 48787, Appl
GENERAL INFORMATION:
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US-60-417-190-48789/c
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US-60-417-190-48788
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SEQ ID NO 48789
LENGTH: 25
TYPE: DNA
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CURRENT FILING DATE: 2002-10-02
NUMBER OF SEO ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEO ID NO 4878
LENGTH: 25
TYPE: DNA
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CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V
SEO ID NO 48787
LENGTH: 25
TYPE: DNA
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Best Local :
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APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
                                                     CURRENT APPLICATION NUMBER: US/60/417,190
CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
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APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
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APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
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Local Similarity 100.0%;
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Pred. No.
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Pred. No.
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US-60-417-190-48792/c : Sequence 48792, Application US/60417190 ; GENERAL INFORMATION:
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US-60-417-190-48791
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US-60-417-190-48790/c
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                                                   RESULT 29
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CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 48791
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 48791, Application US/60417190
GENERAL INFORMATION:
APPLICANT: Giulia Kennedy
APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
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CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 48790
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Best Local Similarity
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Matches 14; Conserv
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APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
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100.0%; Pred. No.
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                                                                                                                                                                                      DB 7; Length 25; le+02;
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APPLICANT: Giulia Kennedy

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APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
CURRENT APPLICATION NUMBER: US/60/417,190
CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 48792
LENGTH: 25
TYPE: DNA
ORGANISM: Homo sapien
US-60-417-190-48792
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; Sequence 48793, Application US/60417190; GENERAL INFORMATION:
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                                                                       ; TYPE: DNA; Homo sapien US-60-417-190-94484
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                                                                                                                NUMBER OF SEQ ID NOS: 122930
SOPTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 94484
LENGTH: 25
TYPE: YELLOW DATE: 2002-10-02
TYPE: YELLOW DATE: 25
Query Match
Best Local S
Matches 14
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CURRENT APPLICATION NUMBER: US/60/417,190
CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 48793
LENGTH: 25
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Best Local Similarity
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APPLICANT: Giulla Kennedy
APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
CURRENT APPLICATION NUMBER: US/60/417,190
CURRENT FILING DATE: 2002-10-02
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APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
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                    Local Similarity
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100.0%; Pred. No.
                    3.8%; Score 14;
100.0%; Pred. No.
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                    DB 7;
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CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 94485
LENGTH: 25
TYPE: DNA
ORGANISM: Homo sapien
Sequence 94487, Application US/60417190

SEQUENCE 94187, Application US/60417190

GENERAL INFORMATION:

APPLICANT: Giulia Kennedy

APPLICANT: Hajime Matsuzaki

APPLICANT: Mei-Mei Shen

TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522

CURRENT APPLICATION NUMBER: US/60/417,190

CURRENT FILING DATE: 2002-10-02
                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/60/417,190
CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 94486
LENGTH: 25
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 94486, Application US/60417190
GENERAL INFORMATION:
APPLICANT: Giulia Kennedy
APPLICANT: Hajime Matsuzaki
APPLICANT: Hei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
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GENERAL INFORMATION:
APPLICANT: Giulia Kennedy
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APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
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Local Similarity 100.0%; Pred. No.
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Local Similarity 100.0%;
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; Sequence 94488, Application US/60417190; GENERAL INFORMATION:
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US-60-417-190-94488
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                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/60/417,190
CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 94489
                                                                   Matches
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                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                     APPLICANT: Giulia Kennedy
APPLICANT: Hajime Matsuzaki
APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
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SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 94487
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APPLICANT: Mei-Mei Shen
TITLE OF INVENTION: Methods for Genotyping Polymorphisms in Humans
FILE REFERENCE: 3522
CURRENT APPLICATION NUMBER: US/60/417,190
CURRENT FILING DATE: 2002-10-02
NUMBER OF SEQ ID NOS: 122930
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
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                                                                                                                                                                    LENGTH: 25
TYPE: DNA
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TYPE: DNA
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TYPE: DNA
                               143 CAGAAAGAAGTGAT 156
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19 CAGAAAGAAGTGAT 6
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                                                                               Local Similarity
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Local Similarity 100.0%;
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100.0%; Pred. No. 1e-
ative 0; Mismatches
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Pred. No.
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; Sequence 19578, Application US/09513999C
; GENERAL INFORMATION:
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US-09-513-999C-19578
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US-09-513-999C-17785
; Sequence 17785, Application US/09513999C
; GENERAL INFORMATION:
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: Sequence 16171, Application US/09513999C
: GENERAL INFORMATION:
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Matches 14; Conserv
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SEQ ID NO 17785
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APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Glordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
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SEQ ID NO 16171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins. FILE REFERENCE: 59.US2.REG CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24
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APPLICANT: Duclert, A.
APPLICANT: Glordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 36681
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Dumas Milne Edwards, J.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/122,487 PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24
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                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: 69
OTHER INFORMATION: h=a or c or t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 100
                                                                                                                                                                                                                                          161 CAAAGTGCCGGGAT 174
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                                                                                                                                                                                                                 43 CAAAGTGCCGGGAT 56
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100.0%; Pred. No. 1.
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Pred. No. 1.1e+02;
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: ORGANISM: Homo sapiens
US-09-513-999C-19578
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                      US-09-513-999C-18596
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NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 19578
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SEQ ID NO 23316
LENGTH: 112
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                                                                                       SOFTWARE: Patent.pm
SEQ ID NO 18596
LENGTH: 116
                                                                                                                                                                                                                                                                                                                        Sequence 18596, Application US/09513999C GENERAL INFORMATION:
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Best Local
Query Match
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CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 199-02-26
NUMBER OF SEQ. ID NOS: 36681
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                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24 PRIOR APPLICATION NUMBER: US 60/122,487 PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                              APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.USZ.REG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins FILE REFERENCE: 59.US2.REG
                                                                                                                                            NUMBER OF SEQ ID NOS: 36681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Homo sapiens
                                                ORGANISM: Homo
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; Pred. No. 1.1e+02;
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Pred. No.
   Score 14;
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 Length 116;
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CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 14544
LENGTH: 125
TYPE: DNA
ORGANISM: Homo sapiens
US-09-513-999C-14544
                                                                                                                                                                                                                                                                                                                   ; NUMBER OF SEQ ID NOS: 36681
; SOFTWARE: Patent.pm
; SEQ ID NO 18527
; LENGTH: 129
; TYPE: DNA
; ORGANISM: HOMO Sapiens
US-09-513-999C-18527
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US-09-513-999C-14544/c
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Sequence 18527, Application US/09513999C
GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
RESULT 44
US-09-513-999C-28218
; Sequence 28218, Application US/09513999C
; GENERAL INFORMATION:
; APPLICANT: Dumas Milne Edwards, J.B.
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Best Local Similarity
Matches 14; Conserv
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Best Local Similarity 100.0%;
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TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
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100.0%; Pred. No. 1.1e+02;
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Pred. No
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APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 25794
LENGTH: 142
TYPE: DNA
ORGANISM: Homo sapiens
US-09-513-999C-25794
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APPLICANT: Glordano, J.Y.
APPLICANT: Glordano, J.Y.
FILE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 28218
LENGTH: 138
TYPE: DNA
ORGANISM: Homo sapiens
US-09-513-999C-28218
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Search completed: November 8, 2002, 02:02:17 Job time: 69.3091 secs
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US-09-513-999C-25794
Sequence 25794, Application US/09513999C
GENERAL INFORMATION:
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Maximum DB seq length: 2000000000
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3: en_

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7: em

9: gt

110: cg

111: cg

111: cg

112: cg

113: cg

114: cg

115: cg

12: cg

12: cg

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

| σ                  | <sub>U</sub>       | 4                  | ω                 | ν                  |                    | Result                       |
|--------------------|--------------------|--------------------|-------------------|--------------------|--------------------|------------------------------|
| 295                | 307                | 314                | 315               | 315                | 315                | score                        |
| 79.9               | 83.2               | 85.1               | 85.4              | 85.4               | 85.4               | Query  Be Match Length DB ID |
| 987                | 747                | 593                | 920               | 707                | 704                | Length                       |
| 14                 | 13                 | 10                 | 9                 | 13                 | 13                 | 8                            |
| BQ668420           | BI917149           | BE386060           | AL538562          | в1913989           | BI919074           | Length DB ID                 |
| BQ668420 AGENCOURT | BI917149 603181571 | BE386060 601276753 | AL538562 AL538562 | BI913989 603180565 | BI919074 603180881 | Description                  |

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## ALIGNMENTS

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| FEATURES                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                   | TITLE                                                                                | AUTHORS                          | REFERENCE          |                                                                                                                                 | ORGANISM     | SOURCE | KEYWORDS | VERSION                | ACCESSION I                | DEFINITION (                                                      | T 1<br>074 |
|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------|--------------------------------------------------------------------------------------|----------------------------------|--------------------|---------------------------------------------------------------------------------------------------------------------------------|--------------|--------|----------|------------------------|----------------------------|-------------------------------------------------------------------|------------|
| Email: cgapbs-remail.nln.gov Tissue Procurement: Life Technologies, Inc. CDNA Library Preparation: Life Technologies, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) CDNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://inage.llnl.gov plate: LLAM11618 row: a column: 21 High quality sequence start: 4 High quality sequence start: 4 High quality sequence start: 4 Location/Qualifiers | Contact: Robert Strausberg, Ph.D. | National Institutes of Health, Mammalian Gene Collection (MGC)<br>Unpublished (1999) | NIH-MGC http://mgc.nci.nih.gov/. | 1 (bases 1 to 704) | Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleoscomi;<br>Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. | Homo sapiens | human. | EST.     | BI919074.1 GI:16200128 | mRNA sequence.<br>B1919074 | 603180881F1 NIH_MGC_121 Homo sapiens cDNA clone IMAGE:5244956 5', |            |

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cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
                                                                                                                                   Contact: Robert Strausberg, Ph.D.
                                                                                                                                                  NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 707)
                                                                                                                                                                                                                                                                                                           BI913989.1 GI:16178286
                                                                                                                                                                                                                                                                                                                                         mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                       BI913989 707 bp mRNA linear EST 16-OCT-20 603180565F1 NIH_MGC_121 Homo sapiens cDNA clone IMAGE:5244569 5',
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/db_xref="taxon:9606"
/clone="IMAGE:5244956"
/clone_lib="NIH_MGC_121"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="Organ: brain; Vector: pCMV-SPORT6; Site_1: Not1; Site_2: EcoRV (destroyed); RNA source anonymous pool of 3 fettal brains, female age 20 weeks, female age 24 weeks, and male age 26 weeks. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.7 kb, insert size range 0.7-3.5 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 017. Note: this is a NIH_MGC Library."
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Pred. No. 1.7e-166;
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435 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 494
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                                                                                                                               Contact: Genoscope
                                                                                                                                                               Full-length cDNA libraries Unpublished (2001)
                                                  Email: segref@genoscope.cns.fr, Web : www.genoscope.cns.fr.
                                                                               BP 191 91006 EVRY cedex - France
                                                                                                       Genoscope - Centre National de Sequencage
                                                                                                                                                                                       \text{Li}, \dot{W}, B., \text{Gruber}, C., \text{Jessee}, J. and \text{Polayes}, D. \text{Full-length} cDNA libraries and normalization
                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/clone="IMAGE:5244569"
/clone_lib="NIH_MGC_121"
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601276753F1 NIH_MGC_20 Homo sapiens
                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                           Unpublished (1999)
                                                                                                                                                                                                NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                          mRNA sequence.
                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov
                                                                                                                                                                                                                                            (bases 1 to 593)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note-"Organ: Fetal brain; Vector: pCMVSPORT 6; 1st strand cDNA was primed with a Noti-cligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and cloned into the Not I and Eco RV sites of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="LTI_F1013_FBrn1"
/dev_stage="pooled tissue from post conception fetuses
week, 24 week and 26 week)"
/lab.host="DH10B"
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/db_xref="taxon:9606"
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100.0%; Pred. No. 1.8e-166;
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Contact: Robert Strausberg, Ph.D.

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Life Technologies, Inc.

cDNA Library Preparation: Life Technologies, Inc.

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                               B1917149 747 bp mRNA linear EST 16-OCT-2001 603181571F1 NIH_MGC_121 Homo sapiens cDNA clone IMAGE:5245648 5',
                                                                                                                                                                       NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                       Homo sapiens
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                    BI917149.1 GI:16181111
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BI917149
                                                                                                                                                       Unpublished (1999)
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                                                                                                                                                                                                                  (bases 1 to 747)
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166 c 183 g 105 t
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/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: skin; Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Organ: skin; Vector: pTTB7; Site_1: XhoI; Site_2:
cloned into EcoRI/XhoI sites using the following 5'
adaptor: GGCACGAG(G)... Size-selected >500bp for average
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/clone_lib="NIH_MGC_20"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        307;
                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                 Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                       NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 987)
                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                BQ668420.1 GI:21778667
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Plate: LLAM11619 row: n column: 17
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                                                                                                                                                                                                                                               , mRNA sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE:5245648"
/clone_lib="NIH_MGC_121"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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Pred. No. 5.7e-162;
                                                                                                                                                                                                                                               987 bp
102 Homo F
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sapiens
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Toshiyuki and Piero Carninci (RIKEN) cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
CDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki
                                                                                                                  Contact: Robert Strausberg, Ph.D.
                                                                                                                                        Unpublished (1999)
                                                                                                                                                      NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                          Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                 mRNA sequence.
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Plate: LLCM2457 row: m column: 10
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DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be
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/tlosue_type="epidermoid carcinoma, cell line"
/tlosue_type="epidermoid carcinoma, cell line"
/lab_host="DH10B (phage-resistant)"
/note="Organ: salivary gland; Vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Site=2: EcoRI; cDNA made by oligo-dT priming the
following 5' adaptor: GGCAGGGG;
Library constructed
by Ling Hong in the laboratory of Gerald M. Rubin
(University of California, Berkeley) using ZAP-CDNA
synthesis kit (Stratagene) and Superscript II RT (Life
Technologies). Note: this is a NIH_MGC Library."
96 a 314 c 285 g 191 t 1 others
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/db_xref="taxon:9606"
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Clone distribution: MGC clone distribution information
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
plate: LLAM11792 row: p column: 10
High quality sequence stop: 690.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                        340 bp mRNA linear EST 21-
y183b07.r1 Soares infant brain 1NIB Homo sapiens cDNA clone
IMAGE:44619 5', mRNA sequence.
                                                                                 Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and Wilson, R.
Washington University School 4444 Forest Park Parkway,
                                           Unpublished (1995)
Contact: Wilson RK
                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 340)
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/db_xref="taxon:9606"
/clone="IMAGE:5312889"
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/tissue_type="hypothalamus"
/lab_host="DH10B"
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Pred. No. 5.5e-149;
School of Medicine way, Box 8501, St.
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This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
Insert Length: 1737 Std Error: 0.00
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Insert Size: 1737
High quality sequence stops: 312
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 1010)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                     Homo sapiens
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/db_xref="GDB:417160"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    CATCAGCTGCTACCACCCCTCTTTGTAAC 301
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          271;
                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                      Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                           NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae.
                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                  BE395206.1
                                                                                                                                                                                                                                                                     BE395206
                                                                                                                                                                                                                                                                                       mRNA sequence.
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Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM9422 row: d column: 05
High quality sequence stop: 649.
Location/Qualifiers
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Tissue Procurement: David N. Louis, M.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
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cDNA Library Preparation: Ling Hong/Rubin Laboratory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /lab_host="DH10B (T1 phage-resistant)"
/note="Organ: brain: Vector: pCMV-SPORT6; Site_1:
Site_2: Sall; Cloned unidirectionally. Primer: Oli
Average insert size 2.3 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
a 344 c 254 g 146 t
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/clone="IMAGE:4153684"
/clone_lib="NCI_CGAP_Brn67"
/tissue_type="anaplastic oligodendroglioma with lp/19q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
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                                                                                           Mammalian Gene Collection (MGC)
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     181 CTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAACGGGCCCAAGGCCCCAAGAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       262 CTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTAACGGGCCAAGGCCCAAGAAAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              82 CAACGACTGCTCCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTG 141
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                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 948)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                  Contact: Robert Strausberg, Ph.D.
                                                                                                                                                        Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                     EST
                                                                                                                                                                                                                                                                                                                                                                                             5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                    BQ882838 948
AGENCOURT_8627686 NIH_MGC_43
CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                       BQ882838.1 GI:22274846
                                                                                                                                                                                                                                                                                                                                                                              BQ882838
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Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
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/clone_lib="NIH_MGC_44"
/tissue_type="endometrium, adenocarcinoma cell line"
/lab_host="Pull0 (phage-resistant)"
/note="Organ: uterus; vector: pOTB7; Site_1: XhoI; Site_2:
ECORI; CDNA made by oligo-dT priming. Directionally
cloned into ECORI/XhoI sites using the following 5'
adaptor: GGCACGAG(G). Library constructed by Ling
in the laboratory of Gerald M. Rubin (University of
California, Beskeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
19 a 192 c 154 g 108 t
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/db_xref="taxon:9606"
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Plate: LLCM2492 row: k column:
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Contact: Robert Strausberg, Ph.D.

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Life Technologies, Inc.

cDNA Library Preparation: Life Technologies, Inc.

cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LINL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:

http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                       NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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               Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 570)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                 Contact: Robert Strausberg, Ph.D.
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Plate: LLAM11424
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/clone="IMAGE:5181585"
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                                    Index (http://www.tigr.org/tdb/hgi/hgi.html)
Seq primer: M13 Reverse.
                                                            For clone availability, additional sequence and expression information related to this EST, please check the TIGR Human Gene
                                                                                                                Email: arkerlav@tigr.org
                                                                                                                                                                             The Institute for Genomic Research 9712 Medical Center Drive, Rockvil
                                                                                                                                                                                                                                          Contact: Kerlavage, AR
                                                                                                                                                                                                                                                             Other_ESTs: EST56848 THC106611
                                                                                                                                                                                                                                                                                                                                   Adams, M.D., Kerlavage, A.R., Fields, C. and Venter, J.C. 3,400 expressed sequence tags identify diversity of transcripts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EST56849 Infant brain Homo sapiens cDNA 5' end, mRNA sequence.
                                                                                                                                                                                                                             Bioinformatics
                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo Sapiens
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                                                                                                                                                                3018699056
                                                                                                                                                                                                                                                                                                                               human brain
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               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note-"Organ: brain; Vector: pCMV-SPORT6; Site_1: Not1; Site_2: ECORV (destroyed); RNA source normal medulla from anonymous male age 27. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.3 kb, insert size range 0.9-3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 013. Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /tissue_type="medulla"
/lab_host="DH10B"
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/clone_lib="NIH_MGC_119"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          290 CCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGGAAGTTCTGCCTCGGCCCTCA 344
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       235;
                                                                                                                                                                                                                                                                                  found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov Plate: LLCM1759 row: e column: 01 High quality sequence stop: 585.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 670)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
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                                                                                                                                                                                                                                                                                                                                                                  CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can
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            /clone="IMAGE: 876392"
/clone=lib="NIH_MGC_43"
/clone=lib="NIH_MGC_43"
/tissue_type="normal pigmented retinal epithelium"
/tissue_type="normal pigmented retinal epithelium"
/tissue_type="normal pigmented retinal epithelium"
/tissue_type="normal pigmented retinal epithelium"
/tissue_type="normal pigmented retinal epithelium"
/tab_host="DH10B (phage-resistant)
/note="Organ: eye; Vector: pOTB7; Site_1: XhoI; Site_2:
ECORI; cDNA made by oligo-dT priming. Directionally
cloned into ECORI/XhoI sites using the following 5;
adaptor: GGCACGAG(G). Library constructed by Ling
nn the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
                                                                                                                                                                                                            /db_xref="taxon:9606"
                                                                                                                                                                                                                             organism="Homo sapiens"
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/db_xref="ARCC (Inhost):150796"
/db_xref="taxon:9606"
/clone_lib="Infant brain"
/sex="female"
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Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov Plate: LLCM760 row: 1 column: 21
High quality sequence stop: 651.
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National Institutes of Health, Mammalian Gene Collection (MGC)
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1 (bases 1 to 654)
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Note: this is a NIH_MGC Library. |"
198 c 165 g 163 t
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/db_xref="taxon:9606"
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Pred. No. 4.3e-115;
Score 205;
Pred. No.
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2.3e-104;
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487
                    209 TGTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTT 268
                                                                             427 GAAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCC
                                                                                                                   149 GAAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCC 208
                                                                                                                                                                              Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N. Louis, M.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Preparation: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC Clone distribution information can be
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National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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BI554034
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 698)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High quality sequence stop:
                                                                                                                                                         53.1%; Score 196; DB 13; ilarity 100.0%; Pred. No. 2.8e-99; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                            /lab_host="DH10B (T1 phage-resistant)"
/note="Organ: brain; Vector: pCMV-SPORT6; Site_1
Site_2: Sall; Cloned unidirectionally. Primer: C
Average insert size 2.3 kb. Constructed by Life
Technologies, Note: this is a NCI_CGAP Library."
                                                                                                                                                                                                                                                                                                                                                                                               /db_xref="taxx0n:9606"
/clone="1mAGE:5309327"
/clone_ltb="NCI_CGAP_Brn67"
/tissue_type="anaplastic oligodendroglioma with lp/19q
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                                                                                                                                                                                                                                                                                              56 ATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTC 115
                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                AGCTGCTGCAACACCCCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCC 334
                                                                                                                                                         TCGGCCCTCA 344
                                                                                                                              TCGGCCCTCA 245
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                                                                                                                                                                                                                                                                                                                                                           190;
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              mRNA sequence.
AA582576
                                        nn55e03.s1 NCI_CGAP_Kid6 Homo sapiens
AA582576.1 GI:2359936
                                                        AA582576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Assessment of gene expression patterns in a model of colon tumor metastasis using a 19,200 element cDNA microarray Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: John Quackenbush
The Institute for Genomic Research
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seq primer: Reverse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9712 Medical Center Dr.,
Tel: 301 838 3528
Fax: 301 838 0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hegde, P., Qi, R., Abernathy, K., Dharap, S., Gaspard, R., Ga, I.E., Saeed, A.I., Sharov, V., Lee, N.H., Yeatman, T.J. and
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ohng@tigr.org
                                                                                                                                                                                                                                                                                                                                                                                                                          /clone_lib="MAGE resequences, MAGC"
/note="Vector: pBluescriptSKm"
150 c 134 g 134 t
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/db_xref="taxon:9606"
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                                                    433 bp
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                                                       mRNA
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                                        NA linear EST 26-SEP-1997 cDNA clone IMAGE:1087804 3',
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Best Local
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KEYWORDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        318 TACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACC
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     NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
National Cancer Institute, Cancer Genome Anat
                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 522)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AI767459 mRNA linear EST 20-DEC-1999 wh26h03.xl NCI_CGAP_Kidll Homo sapiens cDNA clone IMAGE:2381909 3',
                                                                                                                                                                                                                                                          Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mRNA sequence.
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cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-GGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 904 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 407.
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Tissue Procurement: L. Jeffr
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 433)
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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/lab_host="SOLR (kanamycin resistant)"
/lab_host="SOLR (kanamycin resistant)"
/note="Organ: kidney; Vector: Bluescript SK; Site_1:
/note="Organ: kidney; Vector: Bluescript SK; Site_1:
/note="Organ: kidney; Vector: Bluescript SK; Site_1:
/note="Organ: Site_2: XhoI; Cloned unidirectionally. Primer:
/note="Organ: Site_2: XhoI; Cloned unidirectionally. Primer:
/note="Organ: Site_2: XhoI; Cloned unidirectionally. Primer:
/note="Organ: Site_2: XhoI; Cloned unidirectionally. Primer:
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/db_xref="taxon:9606"
/clone="IMAGE:1087804"
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Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                        BM974798 541 bp mRNA linear EST 21-M
UI-CF-EC1-acd-i-10-0-UI.sl UI-CF-EC1 Homo sapiens cDNA clone
UI-CF-EC1-acd-i-10-0-UI 3', mRNA sequence.
                                                        Contact: McCray, PB
                                                                                                                                                           Bonaldo, M.F.,
                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 541)
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                                       McCray Lab
                                                                               97044477
                                                                                                                     discovery
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2024 University of Iowa
                     University of Iowa
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Location/Qualifiers
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                                                                                                Res. 6 (9), 791-806 (1996)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fatima Bonaldo.
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/clone="IMAGE:2381909"
/clone_lib="NCI_CGAP_Kidl1"
/lab_host="DH10B"
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                                                                                                                                                             Lennon, G. and Soares, M.B.
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Med Labs, Iowa City, IA 52242, USA
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FEATURES
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AV717094/c
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                           AUTHORS
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Tissue Procurement: Dr. M. J. Welsh, University of Iowa
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Researchers may obtain clones from Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genetics (www.resgen.com).
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Fax: 319 356 7171
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AV717094 DCB Homo sapiens
AV717094
AV717094.1 GI:10814246
1 (bases 1 to 651)
Xu,X., Gu,J., Liu,F
Song,H., Gu,Y., Yan
                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                          Homo sapiens
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/note="Organ: Lung: Vector: pTTT3-Pac (Pharmacia) with a
modified polylinker; Site_1: EcoR I; Site_2: Not I;
UT-CF-ECI is a normalized cDNA library containing the
following tissue(s): Normal lung from adult and from fetal
day 64, day 87, week 19 and week 42. The library was
constructed according to Bonaldo, Lennon and Soares,
Genome Research, 6:791-806, 1996. First strand cDNA
synthesis was primed with an oligo-dT primer containing a
Not I site. Double stranded cDNA was ligated to an EcoR I
adaptor, digested with Not I, and cloned directionally
into pTTT3-Pac vector. The oligonucleotide used to prime
the synthesis of first-strand cDNA contains a library tag
sequence that is located between the Not I site and the
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/db_xref="taxon:9606"
/clone="UI-CF-EC1-acd-i-10-0-UI"
/clone_lib="UI-CF-EC1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TAG_SEQ=AAGTGCTTAC"
142 c 120 g
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/dev_stage="Adult and Fetal"
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     Liu,F., Qu,J., Zhao,M., Li,Y., Hua
., Yang,Y., Gao,G., Xiao,H., Li,N.,
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                              Huang, Q.,
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            Gao, X.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 175;
       Email: paul-mccray@uiowa.edu
Tissue Procurement: Dr. M. J. Welsh, University of Iowa
cDNA Library preparation: Dr. M. Bento Soares, University of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Researchers may obtain clones from Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                          BM980701 652 bp mRNA linear EST 21-M
UI-CF-EN1-add-j-04-0-UI.Sl UI-CF-EN1 Homo sapiens cDNA clone
UI-CF-EN1-add-j-04-0-UI 3', mRNA sequence.
                                                                                                                                                                               University of Iowa
                                                                                                                                                                                                   McCray Lab
                                                                                                                                                                                                               Contact: McCray, PB
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                                                                                                                                                                                                                                                                     discovery
                                                                                                                                                                                                                                                                                                   1 (bases 1 to 652)
Bonaldo, M.F., Lennor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Zeguang Han
Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong,
T01: 86-21-50801919(ex.45)
                                                                                                                                                                                                                                                                                 Normalization and subtraction: two approaches
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This clone is available at CHGC in Shanghai
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                                                                                                                        University of Iowa Med Labs, Iowa City, IA 52242, USA
319 356 4866
319 356 7171
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(www.resgen.com).
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/lab_host="BM25.8"
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2024 University of Iowa Med Labs, Iowa City, Tel: 319 356 4866
                                           University of Iowa
                                                               McCray Lab
                                                                               Contact: McCray, PB
                                                                                                                                                      discovery
                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 738)
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738 bp mRNA linear UI-CF-EN1-adh-b-10-0-UI.S1 UI-CF-EN1 Homo sapiens UI-CF-EN1-adh-b-10-0-UI 3', mRNA sequence.
                                                                                                           97044477
                                                                                                                                 Genome
                                                                                                                                                                    Normalization and subtraction: two approaches to facilitate gene
                                                                                                                                                                                               Bonaldo, M.F.,
                                                                                                                                                                                                                                                                                      Homo sapiens
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/note-"Organ: Lung; Vector: pT7T3-Pac (Pharmacia) with a
modified polylinker: Site_1: EcoR I; Site_2: Not I;
UI-CF-ENI is a normalized cDNA library containing the
following tissue(s): Primary Lung Cystic Fibrosis
Epithelial Cells. The library was constructed according to
Bonaldo, Lennon and Soares, Genome Research, 6:791-806,
1996. First strand cDNA synthesis was primed with an
oligo-dT primer containing a Not I site. Double stranded
cDNA was ligated to an EcoR I adaptor, digested with Not
I, and cloned directionally into pT7T3-Pac vector. The
oligonucleotide used to prime the synthesis of
first-strand cDNA contains a library tag sequence that is
located between the Not I site and the (dT)18 tall. The
                                                                                                                             Res. 6 (9), 791-806 (1996)
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TAG_LIB-UI-CF-EN1
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/clone="UI-CF-EN1-add-j-04-0-UI"
/clone_lib="UI-CF-EN1"
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                                                                                                                                                                                             Lennon, G. and Soares, M.B.
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Best Local Similarity
                                                                                                                                              ORGANISM
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutherla; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 890)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                                                               mRNA sequence.
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Tissue Procurement: Dr. M. J. Welsh, University of Iowa
cDNA Library preparation: Dr. M. Bento Soares, University of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Researchers may obtain clones from Research
Genetics (www.resgen.com).
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/clone_lib="UI-CF-EN1"
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lone IMAGE:4026252
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               449 DP mRNA linear EST 29-OCT-1998 qe22c09.xl Soares_fetal_lung_NbHL19W Homo sapiens cDNA clone IMAGE:1739728 3', mRNA sequence.
                                                                                                                                                                          Unpublished (1997)
                                                                                                                                                                                              Tumor Gene Index
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 449)
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                                                                                                                                                                                                           NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1999)
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/lab_host="DH10B (phage-resistant)"
/note="norgan: skin; Vector: pOTB7; Site_1: XhoI; Site_2:
ECORI; cDNA made by oligo-dT priming. Directionally
cloned into ECORI/XhoI sites using the following 5'
adaptor: GCGACGAG(G). Size-selected >500bp for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
a 258 c 239 g 161 t
Location/Qualifiers
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/clone_lib="NIH_MGC_20"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGG 336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACC 396
                                                                                                                                                                                                                                                        Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 878)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                      CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov Plate: LLCM856 row: 1 column: 09
                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                       Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                   BF125664.1 GI:10964704
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                601763237F1 NIH_MGC_20 Homo sapiens
                                                                                                                        High quality sequence stop: 598.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
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Pred. No. 2.5e-86;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 878 bp
Vector: pOTB7; Site_1: XhoI; Site_2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mRNA
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Best Local
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                                                                                                                                                                                            source
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                                                                                                                                                                                                                                                                                                                                                      Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1995)
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                                                                                                                                                                                                                          quality sequence stop: 326.
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                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                               . 395
                                                                                                  female'
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RESULT 27 BF125664

BF125664

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290

230

277

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397

ECORI; CDNA made by oligo-dT priming. Directionally cloned into EcORI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)." a 194 c 251 g 155 t

86 GACTGCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAAACGTTCAAGACATGTGTCAG 145 45.3%; 5--100.0%; Pred. No. 100.0%; Pred. No. 100.0%; Pred. No. Score 167; DB 12; Pred. No. 7.3e-83; Length 878; Indels 0, Gaps 0,

300 GACTGCTCCTCCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTCAG 359

BASE COUNT ORIGIN

101

Matches

173;

206 GCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGG 252

GCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGG 466

yg20f09.rl Soares infant brain 1NIB mRNA Нопо linear sapiens o r EST 17-APR-1995 cDNA clone

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo

1 (bases 1 to 395)
Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holma,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterst,R., Williamson,A., Wohldmann,P. and Wilson,R.
The WashU-Merck EST Project Hawkins, M., Holman Waterston

Washington University School of Medicine Louis, MO 63108

Email: est@watson.wustl.edu
Insert Size: 1628
High quality sequence stops: 326 Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Insert Length: 1628
Std Error: 0.00

REFERENCE AUTHORS

TITLE

SOURCE KEYWORDS VERSION ACCESSION DEFINITION

EST

BF125664

ORGANISM

COMMENT

FEATURES

source

/organism="Homo sapiens"

/clone\_lib="Soares infant brain 1NIB

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ACCESSION
VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1810
Fax: 314 286 1810
Email: est@watson.wustl.edu
Insert Size: 1633
High quality sequence stops: 287
Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Insert Length: 1633 Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             High quality sequence stop: 287.
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H17189.1
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429 bp mRNA linear EST 29-,
ym37f07.rl Soares infant brain lNIB Homo sapiens cDNA clone
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/note="Organ: Lafmid BA; Site_1: N.
/note="Organ: Lafmid BA; Site_1: N.
/note="Organ: Lafmid BA; Site_1: N.
/note="Organ: Lafmid BA; Site_1: N.
/note="Organ: Lafmid BA; Site_1: N.
/note="Organ: Lafmid BA; Site_1: N.
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/note="Organ: Lafmid BA; Site_1: N
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            double-stranded
                                                                                                                                                                                                                                      /dev_stage="73 days post natal"
                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:9606"
/clone="IMAGE:50205"
                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="GDB:423015"
                                                                                                                                                                                                                                                                                      /sex="female"
                                                                                                                                                                                                                                                                                                               /clone_lib="Soares infant brain lNIB"
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Pred. No. 2 1e-82;
cDNA was ligated to Hind III adaptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                      cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCT-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 787 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
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AI299246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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              Ω
                       /clone="IMAGE:1900098"
/clone_lib="NCI_CGAP_Kid5"
/tissue_type="2 pooled tumors (clear cell type)"
/lab_host="DH10B"
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127 c 108 g 107 t 5 others
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/db_xref="taxon:9606"
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qd98e07.xl Soares_testis_NHT Homo sapiens cDNA clone
3', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortlum/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 960 Std Error: 0.00
Insert Length: 960 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
National Cancer Institute, Cancer Genome Anat
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                                                                                                                                                                                                                                    constructed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1737540"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="Soares_testis_NHT"
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100.0%; Pred. No. 6.6e-74;
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                                                0;
                                                Score 148; DB 9;
Pred. No. 3.2e-72;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: cgapbs r@mail.nih.gov

The sequence contained an oligo-dT track that was present in the oligonuclectide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonafide poly A tail.cDNA Library Preparation: M.B. Soares Lab Clone distribution: NCI-CGAP clone distribution information can be found through the
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UI-H-BIO-aag-b-04-0-UI.s1 NCI_CGAP_Sub1 Homo sapiens cDNA clone IMAGE:2709078 3', mRNA sequence.
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National Cancer Institute, Cancer Genome Ana
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           /Clone_lb="WCI_CGAP_SUDI"
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/db_xref="taxon:9606"
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/clone_lib="NCI_CGAP_Subl"
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NCI_CGAP_Col0 pool 1 LLAM 2644-2653,
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Insert Length: 941 Std Error: 0.00
Seg primer: -40mi3 fwd. ET from Amersham
High quality sequence stop: 510.
                                                                                                                                                                                                                                                                                                                                                                                            CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
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Unpublished (1997)
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AA776783
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National Cancer Institute, Cancer Genome Anat
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                subtracted library contained 530,000 recombinants.
Subtraction was performed as previously described [Bonaldo, Lennon & Soares (1996): Normalization and Subtraction:
Two Approaches To Facilitate Gene Discovery. Genome
Research 6,791-806.
TAG_LIB-NCI_CGAP_Kid5
Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the
                                                                                                                                                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="1276662"
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                                                                                                                                              /lab_host="DH10B"
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                                                                                                                                                                    /sex="male"
                                                                                                                                                                                       /clone_lib="Soares_testis_NHT"
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al Similarity
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Tissue Procurement: ATCC
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Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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1 (bases 1 to 794)
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                                                                                          /tissue_type="choriocarcinoma"
/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: placenta; vector: pOTB7; Site_1: XhoI;
Site_2: EcoRI; cDNA made by oligo-dT priming.
Directionally cloned into EcoRI/XhoI sites using the
following 5 adaptor: GGCACGAG(G). Size-selected >500bp
for average insert size 1.8kb. Library constructed by
Ling Hong in the laboratory of Gerald M. Rubin (University
of California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
a 227 c 217 g 155 t
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                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:3845440"
/clone_lib="NIH_MGC_21"
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                  38.8%;
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100.0%;
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              Score 143; DB 12;
Pred. No. 2.6e-69;
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Pred. No. 2.3e-69;
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                                 Length 794;
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BE730800
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TITLE
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Best Local
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                                                                                                                                                                  AAGTGATGGAGCAAAGTGCCGGG 426
                                                  AAGTGATGGAGCAAAGTGCCGGG 172
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1 (bases 1 to 1134)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov Tissue Procurement: ATCC
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EST.
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601570755F1 NIH_MGC_21 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                              329
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/db_xrafe"taxon:9606"
/clone_Iib="NIH_MGC_21"
/clone_Iib="NIH_MGC_21"
/tissue_type="choriocarcinoma"
/lab_host="DH10B (phage-resistant)"
/note="Organ: placenta; Vector: pOTB7; Site_1: XhoI;
/note="Organ: placenta; Vector: pOTB7; Site_susing the
pirectionally cloned into EcoRI/XhoI sites using the
following 5' adaptor: GGCAGAGG(G). Size-selected >500bp
for average insert size 1.8kb. Library constructed by
Ling Hong in the laboratory of Gerald M. Rubin (University
of California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
329 a 320 c 308 g 177 t
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                                                                                                                                                                                                                                                                 38.8%;
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                                                                                                                                                                                                                                                                 Score 143; DB 12;
Pred. No. 2.8e-69;
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BG213596
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DEFINITION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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Creation of genome-wide protein expression libraries using random activation of gene expression
activation of gene expression
                                                                                           BG184021
484 bp mRNA linear EST 21-APR-RST2936 Athersys RAGE Library Homo sapiens cDNA, mRNA sequence.
BG184021
BG184021.1 GI:13705708
EST.
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RST33311 Athersys RAGE Library
BG213596
BG213596.1 GI:13735283
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Tel: 216 431 9900
Fax: 216 361 9596
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                          Homo sapiens
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  (bases 1 to 484)
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/cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression
/note="See 'Creation of Genome-wide Protein Expression',
Libraries using Random Activation of Gene Expression',
Nature Biotechnology, in press. Note that even though the
cell type indicated is HT1080, since a random activation
method was used, these sequence tags are not necessarily
expressed in HT1080 under normal circumstances."

93 c 84 g 75 t
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Pred. No. 2.8e-68;
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MEDLINE
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 781)
Harrington,J.J., Sherf,B., Rundlett,S., Jackson,P.D., Perry,R.,
Cain,S., Leventhal,C., Thornton,M., Ramachandran,R., Whittington,J.,
Lerner,L., Costanzo,D., McElligott,K., Boozer,S., Mays,R., Smith
,E., Veloso,N., Kilka,A., Hess,J., Cothren,K., Lo,K., Offenbacher
,J., Danzig,J. and Ducar,M.
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RST12592 Athersys RAGE Library
BG193459
                                                                         Contact: Scott J. Cain Athersys, Inc.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Harrington,J.J., Sherf,B., Rundlett,S., Jackson,P.D., Perry,R., Cain,S., Leventhal,C., Thornton,M., Ramachandran,R., Whittington,J., Lerner,L., Costanzo,D., McElligott,K., Boozer,S., Mays,R., Smith,E., Veloso,N., Kilka,A., Hess,J., Cothren,K., Lo,K., Offenbacher,J., Danzig,J. and Ducar,M.
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Nat. Biotechnol. 19 (5), 440-445 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        J., Danzig,
Creation of
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Carnegie Ave, Cleveland, OH 44115, USA 216 431 9900
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Location/Qualifiers
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216 431 9900
216 361 9596
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/db_xref="taxon:9606"
/db_xref="taxon:9606"
/clone_lib="Athersys RAGE Library"
/cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression
/note="See 'Creation of Genome-wide Protein Expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         38.2%;
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Pred. No.
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s cDNA, mRNA sequence.
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MEDLINE
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1 (bases 1 to 424)

Harrington,J..., Sherf,B., Rundlett,S., Jackson,P.D., Perry,R., Cain,S., Leventhal,C., Thornton,M., Ramachandran,R., Whittington,J., Lerner,L., Costanzo,D., McElligott,K., Boozer,S., Mays,R., Smith,E., Veloso,N., Klika,A., Hess,J., Cothren,K., Lo,K., Offenbacher,J., Danzig,J. and Ducar,M.

Creation of genome-wide protein expression libraries using random
                                                                                                                                                                                                                                                                                                                                           3201 Carnegie Ave, Cleveland, OH 44115, USA
Tel: 216 431 9900
Fax: 216 361 9596
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Scott J. Cain
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High quality sequence stop: 322
Location/Qualifiers
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BG188218.1 GI:13709905
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/db_xref="taxon:9606"
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/cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression / Inbraries using Random Activation of Gene Expression', Nature Biotechnology, in press. Note that even though the cell type indicated is HT1080, since a random activation method was used, these sequence tags are not necessarily expressed in HT1080 under normal circumstances."

87 a 180 c 193 g 221 t
/organism="Homo sapiens"
/db_xref="taxon:9606"
/db_xref="taxon:9606"
/clone_lib="Athersys RAGE Library"
/cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression',
Noter"See 'Creation of Genome-wide Protein Expression',
Nature Biotechnology, in press. Note that even though the
cell type indicated is HT1080, since a random activation
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Location/Qualifiers
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RESULT 40
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MEDLINE
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Best Local Similarity
Matches 140; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE
                                                                             Matches 139;
                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AUTHORS
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 323)

Harrington, J.J., Sherf, B., Rundlett, S., Jackson, P.D., Perry, R., Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, J., Leventhal, C., Thornton, McElligott, K., Boozer, S., Mays, R., Smith, E., Veloso, N., Kilka, A., Hess, J., Cothren, K., Lo, K., Offenbacher J., Danzig, J. and Ducar, M.

Creation of genome-wide protein expression libraries using random activation of gene expression
Nat. Biotechnol. 19 (5), 440-445 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BG192926 323 bp mRNA linear EST 21-APR-2
RST12050 Athersys RAGE Library Homo sapiens cDNA, mRNA sequence.
BG192926
BG192926.1 GI:13714613
                                                                                                                                                                                                                                                                                                                                                                                                                                              Athersys, Inc.
3201 Carnegie Ave, Cleveland,
Tel: 216 431 9900
Fax: 216 361 9596
                                                                                                                                                                                                                                                                                                                                                                                                 Email: scain@athersys.com
High quality sequence stop: 323.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Scott J. Cain
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                                                                           37.7%; So illarity 100.0%; I Conservative 0;
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a 102 c 99 g 101 t 1 others
                                                                                                                                                                Inote-"see 'Creation of Genome-wide Protein Expression Libraries using Random Activation of Gene Expression', Nature Biotechnology, in press. Note that even though the cell type indicated is HT1080, since a random activation method was used, these sequence tags are not necessarily expressed in HT1080 under normal circumstances."
                                                                                                                                                                                                                                                                                                        /clone_lib="Athersys RAGE Library"
/cell_line="HT1080"
                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    37.9%; or 100.0%; Pred. No. - 100.0%; Pred. No. - Mismatches
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Pred. No. 1.1e-67;
                                                                           Score 139; DB 12;
; Pred. No. 3.6e-67;
0; Mismatches 0;
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                                                                                                              Length 323;
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                                                                               0;
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RESULT 42 BG215210 LOCUS DEFINITION

BG215210 324 bp mRNA linear EST 21-APR-2 RST34874 Athersys RAGE Library Homo sapiens cDNA, mRNA sequence.

EST 21-APR-2001

| nes 139; CONSETVACIVE 0;  34 CTTTGCGCTGCAAATCCAGTGCTACCA | //db_xref="caron: 5606" //db_xref="caron: 5606" //db_xref="caron: 5606" //clone_lib="Athersys RAGE //cell_line="HT1080" /note="See 'Creation of G. Libraries using Random Acc Nature Biotechnology, in cell type indicated is HT method was used, these senter the caronic state of G.  method was used, these senter of G.  and S. T. Score 139: atch 37.7%; Score 139: atch 37.7%; Score 139: atch 37.7%; Score 139: atch 37.7%; Score 139: | Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc.  Inc. | Call, S., Leventhal, C., Tho<br>, Lerner, L., Costanzo, D.,<br>,E., Veloso, N., Klika, A.,<br>,J., Danzig, J. and Ducar, M.<br>Creation of genome-wide pr<br>activation of gene express<br>AL Nat. Biotechnol. 19 (5), 4<br>NE 21227151 | ACCESSION BG210457 VERSION BG210457.1 GI:13732144  KEYWORDS EST. SOURCE human. ORGANISM Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Eukaryota; Metazoa; Chordata; Craniata; AUTHORS Harrington, J.J., Sherf, B., Rundlett, S., AUTHORS Harrington, J.J., Sherf, B., Rundlett, S., | GATGGAGCAAAGTGCCGGG 139  GATGGAGCAAAGTGCCGGG 139  m BG210457  N RST30002 Athersys RAGE Library Homo | Db 61 CTCCCCCGAGTTCATTGTGAATTGCAGGTGAACGTTCAAGACATGTGTCAGAAAGTGCCGGG 172  Qy 154 GATGGAGCAAAGTGCCGGG 172 |
|----------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------|
| 993<br>60<br>153                                         | Library"  enome-wide Protein Expritivation of Gene Express Press. Note that even to 1080, since a random acquence tags are not nec normal circumstances."  DB 12; Length 323; ce-67; Length 323;                                                                                                                                                                                                                                              | 4115, USA                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | ies                                                                                                                                                                                                                                     | a; Euteleostomi;<br>ae; Homo.<br>.D., Perry,R.,                                                                                                                                                                                                                                            | mRNA linear EST 21-APR-2001<br>o sapiens cDNA, mRNA sequence.                                       | TCAAGACATGTGTCAGAAAGAAGT 120                                                                             |

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RESULT 43
BG200048/c
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                                                                                       AUTHORS
                                                                                                                                                                         ORGANISM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        34 CTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTGCTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GATGGAGCAAAGTGCCGGG 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTGCTC 60
               1 (bases 1 to 330)
Harrington, J.J., Sherf, B., Rundlett, S., Jackson, P.D., Perry, R., Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, Lerner, L., Costanzo, D., McElligott, K., Boozer, S., Mays, R., Smitl, E., Veloso, N., Klika, A., Hess, J., Cothren, K., Lo, K., Offenbacher
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                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                              BG200048 330 bp mRNA linear EST 21-APR-7
RST19350 Athersys RAGE Library Homo sapiens cDNA, mRNA sequence.
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Creation of genome-wide protein expression libraries using random activation of gene expression
Nat. Biotechnol. 19 (5), 440-445 (2001)
                                                                                                                                                                                                                                      BG200048.1 GI:13721735
                                                                                                                                              Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 216 431 9900
Fax: 216 361 9596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High quality sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: scain@athersys.com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Athersys, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Scott J. Cain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 324)
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BG215210.1 GI:13741231
  Danzig, J. and Ducar, M.
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/cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression
/Interise using Random Activation of Gene Expression',
Nature Biotechnology, in press. Note that even though the
cell type indicated is HT1080, since a random activation
method was used, these sequence tags are not necessarily
expressed in HT1080 under normal circumstances."

83 C 79 g 72 t
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Location/Qualifiers
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/db_xref="taxon:9606"
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Primates;
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                                                                                                                                                  l (bases 1 to 331)

Rarrington, J. J., Sherf, B., Rundlett, S., Jackson, P.D., Perry, R., Cain, S., Leventhal, C., Thornton, M., Ramachandran, R., Whittington, J., Lerner, L., Costanzo, D., McElligott, K., Boozer, S., Mays, R., Smith, E., Veloso, N., Kilka, A., Hess, J., Cothren, K., Lo, K., Offenbacher, J., Danzig, J. and Ducar, M.

Creation of genome-wide protein expression libraries using random activation of gene expression

Nat. Biotechnol. 19 (5), 440-445 (2001)
                                                   3201 Carnegie Ave,
Tel: 216 431 9900
Fax: 216 361 9596
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RST815 At
BG181955
Email: scain@athersys.com
High quality sequence stop: 331.
Location/Qualifiers
                                                                                                          Athersys, Inc.
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21227151
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Tel: 216 431 9900
Fax: 216 361 9596
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Nat. Biotechnol. 19 (5), 440-445 (2001)
                                                                                                                                                                                                                                                                                                                                   Mammalia;
                                                                                                                                                                                                                                                                                                                                                Eukaryota;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: scain@athersys.com
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Athersys RAGE Library Homo
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/cell_line="HT1080"
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/db_xref="taxon:9606"
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Pred. No. 3.7e-67;
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BG204666.1 GI:13726337
EST.
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1 (bases 1 to 331)

Harrington,J.J., Sherf,B., Rundlett,S., Jackson,P.D., Perry,R., Cain,S., Leventhal,C., Thornton,M., Ramachandran,R., Whittington,J., Lerner,L., Costanzo,D., McElligott,K., Boozer,S., Mays,R., Smith,E., Veloso,N., Kilka,A., Hess,J., Cothren,K., Lo,K., Offenbacher,J., Danzig,J. and Ducar,M.
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Nat. Biotechnol. 19 (5), 440-445 (2001)
                                                                                                                                                                                                                                                                  Email: scain@athersys.com
High quality sequence stop: 331.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                          Tel: 216 431 9900
Fax: 216 361 9596
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                                                                                                                                                                                                                                                                                                                                                                                                           Athersys, Inc.
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Libraries using Random Activation of Gene Expression', Nature Biotechnology, in press. Note that even though the cell type indicated is HT1080, since a random activation method was used, these sequence tags are not necessarily expressed in HT1080 under normal circumstances."

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/cell_line="HT1080"
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/db_xref="taxon:9606"
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/cell_line="HT1080"
/note="See 'Creation of Genome-wide Protein Expression
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/db_xref="taxon:9606"
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Best Local Similarity
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Perfect score:
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2054640 seqs, 14551402878 residues
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Pred. No. is the number of results predicted by chance to have

> REFERENCE TITLE JOURNAL AUTHORS

> > Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae 1 (bases 1 to 369)
Xu,J. and Stolk,J.A.
Ovarian tumor sequences and methods of use therefor Patent: WO 0118046-A 199 15-MAR-2001;

Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### Result ACCESSION VERSION KEYWORDS SOURCE ORGANISM RESULT 1 AX093381 LOCUS DEFINITION 00000 O O 0 0 00 145.6 800.2 800.2 37.2 37.2 36.2 36.4 36.2 36.2 284 198.2 159 343.4 336.4 305.2 365.4 365.4 365.4 365.4 365.4 Score AX093381 369 bp Sequence 199 from Patent WO0118046. AX093381 AX093381.1 GI:13509829 Query Match σουσουουουουουουο . 7 7 8 8 8 8 8 129676 209885 147131 136799 190503 19060 150724 180884 177088 2357088 293387 3405 3405 163486 16481 191141 191141 160089 157226 163135 Length DB 65280 179854 189561 39491 40620 164462 169123 199450 204767 204767 2187023 36424 168767 369 946 1524 1524 1832 1890 396 2528 OOOOOOOOOAX362255 AK094501 AX136281 AX093191 AX319944 AB041649 AX136556 AC008018 AC096395 ALT72296 ALT72296 ALB07802 BTEAG1 BTEAG2 AC115890 AC073424 AC014006 AC007813 AC007813 AC0099218 AC099218 AX093381 BC017318 AX358762 AC010974 AC110334 AC128363 AC111928 AP000550 IJ AL390015 AC099562 AC118549 AC007437 AC110739 AC079773 AC124493 LMFL8138 AF239995 AC106546 AC130865 AC124918 AC097561 AC097561 AC120749 SUMMARIES ALIGNMENTS DNA linear Y13430 Bos taurus Y13431 Bos taurus AC115890 Mus muscu AC073424 Homo sapi AC014006 Drosophil AC007813 Drosophil AC007813 Drosophil AC009218 Rattus no AC073438 Homo sapi AC0193562 Homo sapi AC118549 Homo sapi AC118549 Homo sapi AC118549 Homo sapi AC118540 Homo sapi AC118540 Homo sapi AC118540 Homo sapi AC118545 Homo sapi AC118546 Rattus no AC130865 Rattus no AC130865 Rattus no AC12074918 Rattus no AC12074918 Rattus no AC12074918 Rattus no AC120749 Rattus no AC120749 Rattus no AC120749 Rattus no AC120749 Rattus no AC010974 Homo sapi AC110334 Rattus no AC11938 Rattus no AC111928 Rattus no AC006550 Homo sapi AC008018 Homo sapi AC096395 Rattus no AL1772296 Mus muscu AX093381 Sequence BC017318 Homo sapi AX358762 Sequence AX362255 Sequence AK094501 Homo sapi AX136556 Sequence AC07973 Homo sapi AC124493 Mus muscu AC010974 Homo sapi AC010334 Rattus no AC128363 Rattus no AC111928 Rattus no AX136281 Sequence AX093191 Sequence AX319944 Sequence AB041649 Mus muscu Description PAT 30-MAR-2001

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                                                            cDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: National Institutes of Health Intramural
Sequencing Center (NISC),
Shevchenko, Y., Wetherby, K.D., Beckstrom-Sternberg, S.M.,
                                Gaithersburg, Maryland;
Web site: http://www.nisc.nih.gov/
                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                             Submitted (05-NOV-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer
                                                                                                                                                               NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
                                                                                                                                                                                                                                                                     Direct Submission
                                                                                                                                                                                                                                                                                        Strausberg, R
                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 946)
                                                                                                                                                                                                                                                                                                                                                                                                                     946 bp mRNA linear PRI 09-N
Homo sapiens, clone MGC:29643 IMAGE:3641660, mRNA, complete
BC017318
                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                   BC017318.1 GI:16878239
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109 c 90 g
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/db_xref="taxon:9606"
                                                                                                                                                                                                               31 Center Drive, Room 11A03, Bethesda, MD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             location/Qualifiers
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                 nisc_mgc@nhgri.nih.gov
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RESULT 3
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Sequence 15 from Patent W00193983 AX358762
                                          AX358762
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This clone was selected for full length sequencing because it passed the following selection criteria: Hexamer frequency ORE analysis, Similarity but not identity to protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Benjamin,B., Blakesley,R.W., Bouffard,G.G., Brinkley,C., Brooks,S., Dietrich,N.L., Guan,X., Gupta,J., Ho,S.-L., Karlins,E., Legaspi,R., Lim,M., Maduro,Q.L., Mastiello,C., Mastrian,S.D., McCloskey,J.C., McDowell,J., Pearson,R., Snyder,B., Stantripop,S., Thomas,P.J., Tiongson,E.E., Touchman,J.W., Tsurgeon,C., Vogt,J.L., Walker,M.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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/protein_id="AAH17318.1"
/protein_id="AAH17318.1"
/db_xref="G1:16878240"
/translation="MMVLGIAATFCGLFLLPGFALQIQCYQCEEFQLNNDCSSPEFIV
NCTVNVQDMCQKEVMEQSAGIMYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPLCNGPRFKRGGSSASALRPGLATTILFLKLALFSAHC"

292 c 261 g 196 t
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/clone_lib="NIH_MGC_39"
/lab_host="DH10B-R"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                     linear
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                               Baker, K.P., Desnoyers, L., Gerritsen, M.E., Goddard, A., Godowski, P.J., Grimaldi, J.C., Gurney, A.L., Smith, V., Watanabe, C.K. and Wood, W.I.
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Mammalia; Eutheria;
                                                                                                                                                                                                                                        Homo sapiens
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                                       Genentech,
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                                                        Patent: WO 0208288-A 15 31-JAN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AK094501.1 GI:21753575 oligo capping; fis (full insert sequence). Homo sapiens alzheimer cortex cDNA to mRNA, clone_lib:BRALZ2 clone:BRALZ2001350.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AK094501 1832 bp mRNA linear PRI 15-JUL-2002
Homo sapiens cDNA FLJ37182 fis, clone BRALZ2001350, weakly similar
to Homo sapiens Gz-selective GTPase-activating protein (RGS20)
      Isogai,T. and Yamamóto,J.

Direct Submission

Direct Submission

Submitted (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team); 2-6-7

Submitted (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team); 2-6-7

Razusa-Kamatari, Kisarazu, Chiba 292-0812, Japan

Kazusa-Kamatari, Kisarazu, Chiba 292-0812, Japan

Kezusa-Kamatari, Kisarazu, Chiba 392-0812, Japan

Kezusa-Kamatari, Kisarazu, Chiba 392-0815, Fax:81-438-52-3986)

Remail:genomics@hri.co.jp, Tel:81-438-52-3975, Fax:81-438-52-3986)

NEDO human cDNA sequencing project supported by Ministry of Botechnology (RAB); cDNA library

Construction: Helix Research Institute (HRI) (supported by Japan

Construction: Helix Research Institute (HRI) (supported by Japan

Key Technology Center etc.); 5'- & 3'-end one pass sequencing: RAB,
                                                                                                                                                                                                                                                               Kawakami,B., Sugiyama,A., Takemoto,M., Sugiyama,T., Irie,R., Otsuki,T., Sato,H., Wakamatsu,A., Ishii,S., Yamamoto,J., Isono Kawai-Hio,Y., Saito,K., Nishikawa,T., Kimura,K., Yamashita,H., Matsuo,K., Nakamura,Y., Sekine,M., Kikuchi,H., Kanda,K., Wagatsuma,M., Murakawa,K., Kanehori,K., Takahashi-Fujii,A., Oshima,A., Suzuki,Y., Sugano,S., Nagahari,K., Masuho,Y., Nagai, and Isogai,T.
                                                                                                                                                                                                                           Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                              Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Masuho, Y., Nagai, K.

Isono, Y.,

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SOURCE
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                                                           1 (bases 1 to 1890)
Ota,T., Isogai,T., Nishikawa,T., Kawai,Y.,
Hayashi,K.
                                                                                                                                                                                              1890 bp
Sequence 203 from Patent EP1067182.
AX136281
            Patent: EP 1067182-A 203 10-JAN-2001;
Helix Research Institute (JP)
                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                          Secretory protein or membrane protein
                                                                                                                                    Homo sapiens
                                                                                                                                                    human.
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Location/Qualifiers
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/db_xref="G1:21753576"
/db_xref="G1:21753576"
/translation="mwrtcflatafregletlpgfalQIQCYQCEEFQLNNDCSSPEFIV
NCTVNVQDMCQKEVMEQSAGIMYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPLCNGPRPKKRGSSASALRPGLRTTILFLKLALFSAHC"
501 c 513 g 407 t
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/clone_lib="BRALZ2"
/note="cloning vector: pME18SFL3"
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               181 CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTGCATCGCCTCTGCCGGGTACCAGTCCTT 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           354
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA 593
                                                                                                                                                                                                                                                                                                                                                                                         CACCATCCT 662
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                                                                                                           Ovarian tumor sequences and methods of Patent: WO 0118046-A 9 15-MAR-2001; CORIXA CORPORATION (US)
                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1. (bases 1 to 396)
                                                                                                                                                           Xu, J. and Stolk, J.A.
                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                    AX093191.1 GI:13509640
                                                                                                                                                                                                                                                                                  Sequence 9 from Patent W00118046
AX093191
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                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
1. 396
       /note="n = A,T,C or G"
117 c 95 g
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TPLCNGPRPKKRGSSASALRPGLRTTILFLKLASSRHTAELKEMPPPPALFFQPSPPT
PHLPE"
                                                                                                Location/Qualifiers
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274. .771
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Pred. No. 1.1e-95;
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                    Query Match
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Sequence 3 from Patent WO0181634.
AX319944
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                 /protein_id="CAD19382.1"
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/db_xref="GI:17901492"
/db_xref="GI:17901492"
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SYTLSCKLHTFLFEACSYATLLHVLTLSFERVIAICHREQPETSNMSICTNLSSFW
VTSALVALPLLFAMGTEYFLVNVPSHRGITCNRSSTRHAGPETSNMSICTNLSSEST
VTSALVALPLLEAMGTEYFLVNVPSHRGITCNRSGSTRAGGTRPPQLAKSESBESRT
VROSSIFGAFVVYLVVLLSVARWNMAQVLMKSQKGSLAGGTRPPQLAKSESBESRT
ARRQTIIFFLRLIVVTLAVCWMPNQIRRIMAAAKPKHDWTRSYFRAYMILLPFSETFFY
LSSVINPLLYTYSSQOERRVFYQVLCCRLSLOHANHEKRLRVHAHSTTDSAFRYQRPL
LSSVINPLLYTYSSQOERRVFYQVLCCRLSLOHANHEKRLRVHAHSTTDSAFRYQRPL
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LSSVINPLLYTYSSQOERRVFYQVLCCRLSLOHANEKRLRVHAHSTTDSAFRYQRPL
LSSVINPLLYTYSSQOERRVFYQVLCCRLSLOHANEKRLRVHAHSTTDSAFRYQRPL
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/db_xref-"taxon:9606"
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                                                                                                                                 LFASRRQSSARRTEK IFLSTFQSEAEPQSKSQSLSLESLEPNSGAKPANSAAENGFQE
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Pred. No. 2.8e-89;
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    Score 336.4;
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      3.2e-87;
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                                                                     fis (full insert sequence).
Mus musculus (strain:C57BL) adult female cDNA to mRNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus brain cDNA, AB041649
                                                                                                                                                                                                                                                                                                         Submitted (13-APR-2000) Katsuyuki Hashimoto, National Institute of Infectious Diseases, Division of Genetic Resources; 23-1, Toyama 1-chome, Shinjuku-ku, Tokyo 162-8640, Japan (E-mail:khashi@nih.go.jp, URL:http://www.nih.go.jp/yoken/genebank/, Tel:81-3-5285-1111(ex.2120), Fax:81-3-5285-1181)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Osada, N., Kusuda, J., Tanuma, R., Ito, A., Hirata, M., Sugano, S. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Hashimoto,K., Osada,N., Kusuda,J. and Sugano,S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hashimoto, K
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     isolation of full-length cDNA clones from mouse brain
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                                                                                                                                                                                                                                                                  Lab host:
                                                            part of this sequence is reported
                                                                                                                                                                                                                                                                                              http://www.nih.go.jp/yoken/genebank/
/organism="Mus musculus"
/strain="C57BL"
                                          ocation/Qualifiers
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                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 591)
                                                                                                   Secretory protein or membrane protein Patent: EP 1067182-A 478 10-JAN-2001;
                                                                                                                                                Ota, T., Isogai, T., Nishikawa, T., Kawai, Y.,
                                                                                                                                                                                                                    Homo sapiens
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                                                                                         Helix Research Institute (JP)
                                                                                                                                        Hayashi,K.
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                                                                      Location/Qualifiers
                         /db_xref="taxon:9606"
                                        organism-"Homo sapiens"
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/db_xref="GI:7670500"
/tdb_xref="GI:7670500"
/translation="mwtlgiaarfcglfwlpglalQiQcyQceefqlnnDcsspefiv
NCTVNVQDMCQKEVTEQSAGIWYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPLCNGPRPKKRGSSASAIRPGLLTTLLFFHLALCLAHC"
a 487 c 511 g 408 t
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/dev_stage="adult"
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/clone="MNCb-0671"
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                                                                                      Direct Submission

Direct Submission

Submitted (09-JAN-2002) Department of Genetics, Washington

University, 4444 Forest Park Avenue, St. Louis, Missouri 6

On Aug 9, 2001 this sequence version replaced gi:14488388.
                                                                                                                                                                                                                           MO 63108, US
5 (bases 1
                                                                                                                                                                                                                                                        Direct Submission Submitted (09-AUG-2001) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St.
                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
Submitted (10-SEP-2000) Genome Sequencing Center, Washington
University School of Medicine, 4444 Forest Park Parkway, St.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shah, N., Meyer, R., Boyer, E. and Dignan, G. The sequence of Homo sapiens BAC clone RP11-258B17
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Homo Sapiens BAC clone RP11-258B17 from 2, complete sequence
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               Web site: http://genome.wustl.edu/gsc
Contact: sapiens@watson.wustl.edu
------ Summary Statistics
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The RPCI-11 human BAC library was made from the blood of one male donor, as described by Osoegawa,K., Woon,P.Y., Zhao,B., Frengen,E., Tateno,M., Catanese,J.J. and de Jong,P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong and coworkers at the Roswell Park Cancer Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >- 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The clone sequenced to the left is RP11-1L22; the clone sequenced to the right is RP11-159N20, 2000 bp overlap. Actual start of this clone is at base position 1 of RP11-258B17; actual end is at base position 60003 of RP11-159N20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mapping information for this clone was provided by Dr. John D. McPherson, Department of Genetics, Washington University, St. Louis MO. For additional information about the map position of this sequence, see http://genome.wustl.edu/gsc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RP11-258B17 contains a single plasmid region from 1230 to 1239.
Location/Qualifiers
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1397. .3331
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16810.
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16629
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16269
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15532.
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11961
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/db_xref="taxon:9606"
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21300.
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27048. .2
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20618
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29727. .3
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27309
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           /rpt_
34079
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31494
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30738
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27698
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27403.
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18434. .18
                               /rpt_family="(GA)n"
32667. .32787
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32053
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27926. .2
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26656. .2704
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25485..25
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/rpt_family="L2"
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3. .21796
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ORGANISM
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Sequencing vector: M13; 08
Sequencing vector: plasmid; 1008
Chemistry: Dye-primer ET; 08 of reads
Chemistry: Dye-terminator Big Dye; 1008 of reads
Assembly program: Phrap; version 0.990319
Consensus quality: 208484 bases at least Q40
                                                                                                                                                                                                                                                                                                                                                  Submitted (05-JUL-2002) Genome Sequencing Center, 4444 Forest Park
Parkway, St. Louis, MO 63108, USA
On Jul 5, 2002 this sequence version replaced gi:21426614.
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McPherson, J.D. and Waterston, R.H.
Direct Submission
                                                                                                                                                                                                                         Web site:http://genome.wustl.edu/gsc/index.shtml Contact: submissions@watson.wustl.edu
                                                                                                                                                                                                                                                                    Center: Washington University Genome Sequencing Center Center code: WUGSC
                                                                                                                                                                                Center project name: M_BA0462P13
                                                                                                                                                                                                                                                                                                                                                                                                                           Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Parkway, St. Louis, MO 63108, USA 3 (bases 1 to 209885)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (14-JUN-2002) Genome Sequencing Center, 4444 Forest Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The sequence of Mus musculus clone
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McPherson, J.D. and Waterston, R.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                    --- Summary Statistics -----
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98.0%;
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                                                                                                                                                                                                                             TGGCTCCGCACCACCATCCT 369
                                         147131 bp
Homo sapiens BAC clone RP11-159N20
AC010974
AC010974.9 G1:19033964
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Consensus quality: 208911 bases at least Q30 Consensus quality: 209285 bases at least Q20 Insert size: 206000; agarose-fp Insert size: 212112; sum-of-contigs Quality coverage: 0.00 in Q20 bases; agarose-fp Quality coverage: 11.62 in Q20 bases; sum-of-contigs
                                    HTG
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     * NOTE: This is a 'working draft' sequence. It currently consists of 5 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence
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49555. 107384
/note="assembly_name:Contig11"
107485. 209885
/note="assembly_name:Contig12"
44784 c 46164 g 58745 t
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28718. .49454
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11432. .28617
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/chromosome="UNK"
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11431: gap of unknown length
28617: contig of 17186 bp in length
28717: gap of unknown length
49454: contig of 20737 bp in length
49554: gap of unknown length
107384: contig of 57830 bp in length
107484: gap of unknown length
209885: contig of 102401 bp in length
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5 (bases 1 to 147131)
Waterston, R.
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The sequence of Home sapiens BAC clone RP11-159N20
Unpublished (2001)
3 (bases 1 to 147131)
                                                                                                                                                                                                                                                                              Submitted (01-MAR-2002) Department of Genetics, Washington University, 4444 Forest Park Avenue, St. Louis, Missouri 63108, On Mar 1, 2002 this sequence version replaced g1:13435273.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 147131) Sulston, J. E. and Waterston, R.
NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Direct Submission
Submitted (23-MAR-2001) Genome Sequencing Center, Washington
University School of Medicine, 4444 Forest Park Parkway, St.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Waterston, R.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (28-SEP-1999) Genome Sequencing Center, Washington University School of Medicine, 4444 Forest Park Parkway, St.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Toward a complete human genome sequence Genome Res. 8 (11), 1097-1108 (1998)
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(bases 1 to 147131)
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                                                                                                                                                               Contact: sapiens@watson.wustl.edu
                                                                                                                                       Center project name: H_NH0159N20
                                                                                                                                                                                                                                  Center: Washington University Genome Sequencing Center Center code: WUGSC
                                                                                                                                                                                                             Web site: http://genome.wustl.edu/gsc
                                                                                                                                                                                                                                                                                                                                                                            Submission
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This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by restriction digest.

### MAPPING INFORMATION:

Mapping information for this clone was provided by Dr. John D. McPherson, Department of Genetics, Washington University, St. MO. For additional information about the map position of this sequence, see http://genome.wustl.edu/gsc D. Louis

# SOURCE INFORMATION:

Research Genetics, Inc. (http://www.resgen.com) or Pieter de and coworkers at http://www.chori.org
VECTOR: pBACe3.6 The RPCI-11 human BAC library was made from the blood of one donor, as described by Oscegawa, K., Woon, P. K., Zhao, B., Freng Tateno, M., Catanese, J. J. and de Jong, P. J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either Frengen, E. male

NEIGHBORING SEQUENCE INFORMATION:
The clone sequenced to the right is RP11-458A7, 200 bp overlap; the clone sequenced to the right is RP11-258B17. Actual start of this clone sequenced to the right is RP11-159N20; actual end is at base clone is at base position 1 of RP11-159N20; actual end is at base position 147131 of RP11-159N20

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FEATURES
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/note="match to EST BG184021

9743. 9895
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8373. .8683
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984. .1176
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            /note="match to EST 15183. .15321
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118. .1013
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/db_xref="taxon:9606"
/chromosome="2"
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/note="match to EST
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15183. .15321
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13456. .13776
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BG753617 (NID:g14064270)"
                             BG704350 (NID:g13977603)"
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AC110334
AC110334.3 GI:21738206
HTG; HTGS_PHASE1.
                                   Rattus norvegicus clone CH230-140B23, ***, 69 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="match to EST BG200485 (NID:gl3722172)" 19926. .20089
                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="similar to Homo sapiens EST BE963800
(NID:g11767217)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="similar to Homo sapiens EST R54936 (NID:g819192)
yj75b04.s1"
18938. .19409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="similar to Bos taurus EST BE750540 (NID:g10164532)"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="match to EST BG184021 (NID:g13705708)"
15184. .15320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="match to EST H06756 (NID:g870288) y183b07.r1"
15184. .15321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        note="similar to Bos taurus EST BE682268 (NID:g10067956)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'note="similar to Homo sapiens EST AA779875 (NID:g2839206)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          note-"match to EST BF940247 (NID:g12357567)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18637
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note match to EST BE386060 (NID:g9331425)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note="match to EST BG753617 (NID:g14064270)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 note="match to EST BF347279"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="match to EST BG704350 (NID:g13977603)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="match to EST BG200048 (NID:gl3721735)"
L5767. .15789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="match to EST BF347279
15184. .15321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'note="match to EST R54748 (NID:g819270) yj75b04.r1'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rpt_family="(TCCC)n"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note="match to EST H06756 (NID:g870288) y183b07.r1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  note="match to EST BG187166 (NID:g13708853)"
                                                                                                                                                                                                                                                                                                                                                                                      39.5%; Score 145.6; DB 9 96.7%; Pred. No. 1.6e-31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           _family="AT_rich"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   family="AT_rich"
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Rattus norvegicus
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Harris, C., Harris, K., Harris, K., Harris, C., Harris, C., Harris, K., Harris, K., Harris, K., Harris, K., Harris, K., Harris, Holloway, C., Hollins, B., Homsi, F., Howard, S., Huber, J., Hulyk, S., Hume, J., Jackson, L.E., Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S., Kratovic, J., Kureshi, A., Landry, N., Leal, B., Lewis, L.C., Liu, J., Liu, W., Lucier, A., Lucier, R., Luna, R., Ma, J., Kovar, C., Karlsson, E., Lichtarge, O., Lieu, C., Liu, J., Liu, W., Loulseged, H., Lozado, R.J., Lu, X., Lucier, A., Lucier, R., Luna, R., Ma, J., Massey, E., Mathiney, E., McLeod, M. P., Meddor, M., Mertinez, E., Miner, G., Miner, Z., Mitchell, T., Mohabbat, K., Morgan, M., Morris, S., Moser, M., Neal, D., Newtson, J., Newtson, N., Nguyen, A., Nguyen, N., Okwuonu, G., Oragunye, N., Oviedo, R., Pace, A., Payton, B., Peery, J., Perez, L., Pickens, R., Primus, E., Pu, L. L., Quiles, M., Ren, Y., Rives, M., Rojas, A., Rojubokan, I., Rolfe, M., Ruiz, S., Savery, G., Scherer, S., Scott, G., Shen, H., Shooshtari, N., Sisson, I., Sutton, A., Svatek, A., Tabor, P., Tamerisa, A., Tamerisa, N., Thomas, N., Thomas, N., Wang, S., Ward-Moore, S., Warren, R., Washington, C., Watlilams, G., Willamson, A., Wleczyk, R., Wooden, S., Watlitans, G., Wullasson, A., Weczyk, R., Wooden, S., Watlitans, G., Morley, K., Washistock, G. and Gibbs, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (13-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 12, 2002 this sequence version replaced gi:18847022.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (11-FEB-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
                                                                                                                                                                                                                                                                 Center project name: GQLL Center clone name: CH230-140B23
                                                                                                                                                                                                                                                                                                                                                                                                        Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Web site: http://www.hgsc.bcm.tmc.edu/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Center code: BCM
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NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 69 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
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Gaps

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REFERENCE AUTHORS

1 (bases 1 to 190503)

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;

Rattus

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Douthwaite, K. J., Draper, H., Dugan-Rocha, S., Durbin, K.J.,

Barnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M.,

Falls, T., Ferraguto, D., Flagg, N., Ford, J., Foster, P., Erantz, P.,

Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R.,

Gorrell, J.H., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K.,

Harris, C., Harris, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J.,

Hernandez, O., Hodgson, A., Hogues, M., Holloway, C., Hollins, B.,

Homsi, F., Howard, S., Huber, J., Hulyk, S., Hume, J., Jackson, L.E.,

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Karlsson, E., Kelly, S., Khan, U., King, L., Korvah, J., Kovar, C.,

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Li, J., Li, R., Lucier, A., Lucier, R., Lun, R., Ma, J.,

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Massey, E., Mawhiney, E., McLeod, M.P., Meador, M., Mei, G., Metzker, M.,

Massey, E., Mawhiney, E., McLeod, M.P., Meador, M., May, J.,

Peters, L., Pickens, R., Primus, B., Pu, L.L., Qulies, M., Ren, Y.,

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Rives, M., Rojas, A., Rojubokan, I., Rolfe, M., Ruiz, S., Savery, G.,

Scherer, S., Scott, G., Shen, H., Stonek, H., Stone, H.,

Sodergren, E., Sonaike, T., Sparks, A., Stanley, H., Stone, H.,

Sutton, A., Svatek, A., Tabor, P., Tamerisa, A., Tamerisa, K., Tang, H.,

Tansey, J., Taylor, C., Taylor, T., Telfrod, B., Thomas, N., Thomas, S.,

Williams, G., Williamson, A., Wilczyk, R., Wooden, S., Warlington, S.,

Williams, G., Williamson, A., Wilczyk, R., Wooden, S., Warlington, S.,

Williams, G., Williamson, A., Wilczyk, R., Wooden, S., Warling, S.,

Williams, G., Williamson, A., Wilczyk, R., Welson, D.,

Williams, G., Williamson, A., Williamson, A., Wilson, B.,

Williams, G., Williamson, A., Williamson, S., Woolen, S.,

Williams, M., Williamson, A., Williamson, M., Williamson, B.,

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                                                                                                                                                                                                                                                                                                                                              Submitted (19-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wu,C., Wu,Y., Wu,Y.F., Zhou,J., Zorrilla,S., Nelson,D., Weinstock,G. and Gibbs,R.
NOTE: Estimated insert size may differ from sequence length (see http://www.hgysc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 69 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as
                                                                                                   Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 131119 bases at least 040
Consensus quality: 136808 bases at least 030
Consensus quality: 141839 bases at least Q20
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| 21<br>21<br>21<br>22<br>22                                                                                        | 220                           | 0000 000000000000000000000000000000000                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     | 22222222224442222244444222                                                                                                                                                       |
| AAQ60261<br>AAS60261<br>AAC51550<br>AAC35371<br>ABN99125<br>AAK57408                                              |                               | AAF94044 ABT03283 ABL40351 ABQ54231 ABL1933 ABL19930 AAC75880 AAC75880 AAC87591 AAL59341 AAL59341 AAL59363 ABAL6515 AAH64976 AAH64976                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      | ABI'03284 ABI40352 AAF22400 AAF64188 AAF94818 AAF94818 ABI'03085 ABI03282 ABI48768 ABI03282 ABI40350 AAD18690                                                                    |
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Isolated polypeptides associated with ovarian carcinomas, and the

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Best Local S
Matches 369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention provides a number of coding sequences and proteins, the over-expression of which is associated with ovarian carcinoma/cancer. These can be used in the diagnosis, treatment and prevention of ovarian cancer, optionally by gene therapy or in the form of a vaccine. The present sequence is an example of one of these sequences.
                                14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic acids that encode them, useful for the prevention diagnosis and treatment of ovarian cancers - % \left( 1\right) =\left\{ 1\right\} 
                                                                                                     23-MAY-2002
                                                                                                                           WO200239885-A2
                                                                                                                                                                       cytostatic; gene;
                                                                                                                                                                                   Human; ovarian cancer;
                                                                                                                                                                                                       Human ovarian carcinoma associated coding sequence SEQ ID NO: 199
                                                                                                                                                                                                                                 05-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 369 BP; 82 A; 109 C; 90 G; 86 T; 2 other;
                                                                             13-NOV-2001; 2001WO-US45395
                                                                                                                                                                                                                                                                             ABT03274 standard;
           (CORI-) CORIXA CORP
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                                                                                                                                                                                                                                                                             CDNA;
                                                                                                                                                                                                                                 entry)
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                                                                                                                                                                                   ovarian carcinoma;
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Pred. No. 1.8e-110;
0; Mismatches 0;
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                                                                                                           Ovarian
                                                                                                                                                  Ovarian carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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10-SEP-1999;
                     03-APR-2001;
                                            10-JAN-2002
                                                                   US2002004491-A1
                                                                                        Homo sapiens
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                                                                                                                                                                                              ABL48956;
                                                                                                                                                                                                                     ABL48956 standard;
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                                                                                                                        cancer; cancer therapy; vaccine; gene therapy; tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                      2001US-0825294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 82
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99US-0394374
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to ovarian tumour polynucleotides and polypeptides CC that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polypucleotides of the invention are CC useful for detecting a cancer in a pattent, for stimulating and/or CC expanding T-cells specific for a tumour protein, and for inhibiting the CC development of a cancer in a patient. They are also useful for CC stimulating an immune response in a patient, and for treating a cancer in CC a patient and for determining the presence of a cancer in a patient. CC The isolated polynucleotides of the invention are useful for their CC ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and CC preparing ribozyme molecules for inhibiting expression of tumour CC polypeptides in tumour cells. Polypeptides and polynucleotides of the CC invention are also useful in recombinant DNA molecules to direct CC expression of a polypeptide in appropriate host cells. The sequences CC given in records ABL48760-ABL48956 represent polynucleotides encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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15-AUG-2000;
07-SEP-2000;
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(STOL/)
(ALGA/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ovarian carcinoma proteins.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ovarian tumour polypeptide and polynucleotide useful in diagnosis, prevention and/or treatment of cancer, especially ovarian cancer
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361
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                                                                                                                                                                                                                                                         2002-171027/22.
                              CACCATCCT
                                                                                                                                                                                                           CCGCAAGTCCTGTGCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGGTACCAGTCCTT
                                                                                                                                                                                                                                                                                                                         GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC 120
CACCATCCT
                                                               CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCAC
                                                                                            CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCAC
                                                                                                                              CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA
                                                                                                                                                           CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA
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) STOLK J A.
) ALGATE P A.
) FLING S P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page
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2000US-0640173.
2000US-0656668.
2000US-0713550.
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                              369
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Pred. No. 1.8e-110
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RESULT

applications in molecular biology,

including use as hybridisation probes

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ABK33543
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01-DEC-2000;
20-DEC-2000;
20-DEC-2000;
20-DEC-2000;
28-FEB-2001;
10-MAY-2001;
25-MAY-2001;
     The invention relates to one hundred and twenty two nucleic acids encoding PRO polypeptides. The sequences of the 122 PRO polypucleotides encode human secreted proteins. The PRO nucleic acids, polypeptides, agonists and antagonists are useful for treating a PRO related disorder. The PRO polypeptides are useful for diagnosing tumours, especially lung cancer, colon cancer, breast tumour, prostate tumour, rectal tumour or liver tumour. The PRO polypeptides are useful for stimulating the proliferation of, or gene expression, in pericyte cells, for stimulating the proliferation of differentiation of chondrocyte cells, for stimulating the release of tumour necrosis factor-alpha from human blood, for stimulating or inhibiting the proliferation of normal human dermal fibroblast cells. The PRO polypeptide may also be used as molecular weight markers and for tissue typing. The PRO nucleic acids have
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             breast cancer; prostate tumour; rectal tumour; liver tumour; pericyte cell proliferation; chondrocyte cell proliferation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-JAN-2002
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                                                                                                                                                                                                               Claim 2; Figure 15; 359pp; English.
                                                                                                                                                                                                                                                       One hundred and twenty two nucleic acids encoding PRO polypeptides, useful for treating a PRO related disorder and for diagnosing tumours such as lung cancer, colon cancer, breast tumour, prostate tumour, rectal
                                                                                                                                                                                                                                                                                                                                                           Grimaldi JC,
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                                                                                                                                                                                                                                                                                                                                                                                                  (GETH ) GENENTECH INC
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2000WO-US30873.
2000US-253646P.
2000WO-US32678.
2000US-0747259.
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2001WO-US17092
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2001WO-US06520
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2000WO-US20710.
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Smith V, Stephan JF,
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TF. Watanabe CK,
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RESULT 5
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     Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                                  WPI;
                                                                       Χu J,
                                                                                                               14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
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                                                                                                                                                                                                 WO200239885-A2
                                                                                                                                                                                                                                                   Human; ovarian cancer;
                                                                                                                                                                                                                                                                     Human ovarian carcinoma associated coding sequence SEQ ID NO: 205
                                                                                                                                                                                                                                                                                             05-SEP-2002
                                                                                                                                                                                                                                                                                                                                     ABT03277 standard; cDNA; 1619
                                                                                           (CORI-) CORIXA CORP.
                                                                                                                                                                                                                     Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in chromosome and gene mapping. \mathtt{ABK33536}\text{-}\mathtt{ABK33657} represent human protein coding sequences of the invention.
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                                                  2002-500186/53
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                                                                     Stolk JA,
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56; Conservative
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                                                                       Algate
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99.2%;
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Pred. No. 1.1e-109;
0; Mismatches 3;
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RESULT 6
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Best Local Similarity
                                                                                                                                             14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
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  WPI; 2002-500186/53
                                                                                                                                                                                                                                           13-NOV-2001; 2001WO-US45395
                                                                                                                                                                                                                                                                                          23-MAY-2002
                                                                                                                                                                                                                                                                                                                                         WO200239885-A2
                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                               cytostatic; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; ovarian cancer;
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                                                                                              (CORI-) CORIXA CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                 Algate
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                                               Fling
                                                                                                                                                                                                                                                                                                                                                                                                                                                        carcinoma; gene therapy;
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01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
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                                                                                                                                                                                                                                    US2002004491-A1
                                                                                                                                                                                                                                                                                                                                                           Ovarian cancer;
                                                                                                                                                                                                                                                                                                                                                                                                        Ovarian carcinoma sequence isolate 57887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABL40345 standard; cDNA; 1619
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian cancer
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2000US-0561778.
2000US-0640173.
2000US-0656668.
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Best Local Similarity
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(STOL/)
(ALGA/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (STOL/) STOLK J A. (ALGA/) ALGATE P A (FLIN/) FLING S P.
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1619 BP; 399 A; 415 C; 432 G; 373 T; 0 other;
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                             CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA
                                                                CCGCAAGTCCTGTGCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT
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                                                                                                                                                                                                                                                                                                                                                      Conservative
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01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
                                                                                                                                                                                                                                                                                                                        a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the ovarian carcinoma 0591S nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to ovarian tumour polynucleotides and polypeptides that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or useful for detecting a cancer in a patient.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer
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                                                                                                                                                                            GGCAACTTTTTGCGGATTGTTCTTGCTTNCAGGCTTTGCGCTGCAAATCCAGTGCTACCA
                                GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA
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                                                                GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA 180
                                                                                               GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC
                                                                                                                              GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC
                                                                                                                                                                GGCAACTTTTTGCGGATTGTTCTTGCTTCCAGGCTTTGCGCTGCAAATCCAGTGCTACCA
                                                                                                                                                                                                                               366;
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2000US-0656668.
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99.2%;
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                                                                                                                                                                                                                                              Score 365.4; DB 24; Pred. No. 1.2e-109;
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RESULT 9
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11-JAN-2000;
02-MAY-2000;
protein/membrane protein expression. The nucleic acids and complementary sequences may also be used as DNA probes in diagnostic assays (e.g. polymerase chain reactions (PCR)) to detect and quantitate the presence of similar nucleic acid sequences in samples. They may also be used to study the expression and function of secretory proteins/membrane polypeptides and their role in metabolism. The polypeptides may be used as antigens in the production of antibodies against them and in assays to identify modulators (agonists and antagonists) of expression and activity. The antibodies and antagonists may also be used as therapeutic agents to down regulate expression and activity. The antibodies may also
                                                                                                                                                                  This invention relates to nucleic acid sequences AAF93744 - AAF93916 which encode human secretory or membrane proteins represented by AAB88317 - AAB88419. Included in the invention are primers AAF93917 - AAF94295 and AAF62237 - AAF62235 which are used to isolate to CDNA sequences of the invention. The invention also includes methods for the production of antibodies directed against the proteins, and cDNA sequences, which can be used in vaccines. The polynucleotide sequences can be used in gene therapy. The polynucleotide sequences and the proteins they encode may be used in the prevention, treatment and diagnosis of diseases associated with inappropriate secretory.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; secretory protein; membrane protein; vaccine; gene therapy; rheumatoid arthritis; diabetes; ss.
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                                                                                                                                                                                                                                                                                                                                                                  Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                  gene
                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acids encoding secretory proteins/membrane
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                                                                                                                                                                                                                                                                                                                                                                  SEQ ID 203; 609pp + CD ROM; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000JP-0118775
2000JP-0183766
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rug development -
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                                                                                                14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966.
Novel ovarian cancer polypeptide and polynucleotide, useful detecting the presence of ovarian cancer in a patient, and j
                                    WPI; 2002-500186/53.
                                                           χu J,
                                                                                                                                                                                                                                  Human; ovarian cancer; ovarian carcinoma; gene therapy; immunotherapy;
                                                                                                                                                                                                                                                      Human ovarian carcinoma associated coding sequence SEQ ID NO:
                                                                                                                                                                 23-MAY-2002.
                                                                                                                                                                                      WO200239885-A2
                                                                                                                                                                                                                              cytostatic; gene;
                                                                              (CORI-) CORIXA CORP
                                                                                                                                            13-NOV-2001; 2001WO-US45395
                                                                                                                                                                                                                                                                                05-SEP-2002 (first entry)
                                                                                                                                                                                                                                                                                                       ABT03284;
                                                                                                                                                                                                                                                                                                                          ABT03284 standard; cDNA; 1897
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1890 BP; 419 A; 528 C; 533 G; 410 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  be used as diagnostic agents for detecting the presence of the polypeptides in samples (e.g. by enzyme linked immunosorbant assay (ELISA). Examples of diseases which may be treated include rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              arthritis and diabetes.
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                                                         Stolk JA,
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ilarity 99.2%;
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0; Mismatches 3;
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                                                      Molesh DA;
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10-SEP-1999; 99US-0394374
01-MAY-2000; 2000US-0561778
                                                                                                                                                                                                                                                                                                   CDS
                                                                            03-APR-2001; 2001US-0825294
                                                                                                                                                                                       US2002004491-A1.
                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ovarian cancer; cancer therapy; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ovarian carcinoma 01034C/0591S consensus nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-JUN-2002
                                                                                                                                       10-JAN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              580
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                                                                                                                                                                                                                                                                                                Location/Qualifiers 260..685
                                                                                                                                                                                                                                          /product-
                                                                                                                                                                                                                                                                    /*tag=
                                                                                                                                                                                                                                  "Ovarian carcinoma protein O1034C/O591S"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
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Pred. No. 1.2e-109;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene therapy; tumour; cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating and for determining the presence of a cancer in a patient and for treating a cancer in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-AUG-2000;
07-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1897 BP; 435 A; 521 C; 532 G; 407 T; 2 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence represents the ovarian carcinoma 01034C/0591S consensus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current
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                                                                                                                                                                                                                                                                                                GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGGATCATGTA 180
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   CACCATCCT
                                    CACCATCCT 369
                                                                                                   CGGGCCAAGGCCCAAGAAAAGGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCAC
                                                                                                                                    CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTAA
                                                                                                                                                     CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA 300
                                                                                                                                                                                                   CCGCAAGTCCTGTGCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT
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                                                                  CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCAGGGCCAGGGCTCCGCAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                366;
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) ALGATE P A.
) FLING S P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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; 2000US-0656668.
; 2000US-0713550.
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99.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polynucleotide useful in diagnosis, cancer, especially ovarian cancer
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RESULT 12

Query Match Best Local Similarity

95.5**%**; 98.6**%**;

Score Pred.

352.4; DB 21; No. 2.4e-105;

Length 1953;

Sequence 1953 BP; 428 A; 540 C;

524 G;

458 T; 3 other;

o f

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AAF22400/c ID AAF22400 standard; cDNA; 1953 BP.
                                                                                                                                                                                                                                             cc AAF22373 to AAF22421 encode the human secreted proteins given in AAB63134 ct AAB63182. AAB63183 to AAB63231 represent more human secreted proteins can depolypeptides homologous to them. Human secreted proteins have cativities based on the tissues and cells the genes are expressed in. Examples of activities include: immunosuppressive; antiarthritic: cc antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cc erebroprotective; noctropic; neuroprotective; antibacterial; virucide; fungicide; and ophthalmological. The polynucleotides and proteins can be cused to prevent, treat or ameliorate a medical condition in e.g. humans, cm increased in diagnosing a pathological condition or susceptibility to a cardiacy cation. Disorders which are diagnosed or treated include autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative condisorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiacarrest, cerebrovascular disorders e.g. Alzheimar's cerebral condision. Alzheimar's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; secreted protein; diagnosis; immunosuppressive; antiarthritic; antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; ophthalmological; gene therapy; autoimmune disease; neoplasm; rheumatoid arthritis; hyperproliferative disorder; cardiac arrest; cardiovascular disorder; cerebrovascular disorder; cerebral ischaemia; angiogenesis; nervous system disorder; Alzheimer's disease; infection; angiogenesis; nervous system disorder; Alzheimer's disease; infection;
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                                                                                                 ischaemia, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin aging due to subburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and i chemotaxis. The polypeptides can also be used as a food additive or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 440; 533pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isolated nucleic acid molecule encoding a human secreted protein
used in preventing, treating or ameliorating a medical condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (HUMA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-APR-1999; 99US-0128694.
20-JAN-2000; 2000US-0176931.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           used in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200061629-A1
                                                         preservative to increase or decrease storage capabilities. AAF22364 to AAF22372 and AAB63133 represent sequences used in the exemplification (
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ROSEN C A.
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                                            invention.
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        Claim 1; Page 443-444;
                                 preventing, treating or ameliorating a disorder, Parkinson's diseases and cancers -
                                         Nucleic acid molecules encoding human secreted proteins, used in preventing, treating or ameliorating a disorder, e.g. Alzheimer's
                                                                                    P-PSDB;
                                                                                                                         Rosen CA,
                                                                                                                                                                                        11-JUN-1999;
                                                                                                                                                                                                                01-JUN-2000;
                                                                                                                                                                                                                                         21-DEC-2000
                                                                                                                                                                                                                                                                  WO200077026-A1
                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                   Human secreted protein gene 13 SEQ ID NO:23
                                                                                                                                                                                                                                                                                                                                                                                                                                             06-APR-2001 (first entry)
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                                                                                    2001-071258/08
DB; AAB75518.
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                                                                                                                                                 ROSEN
                                                                                                                         Ruben SM,
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                                                                                                                                                                                      99US-0138630.
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        542pp; English
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                                                                                                                                                                                                                                                                                                                            disorder; skin ageing; chemotaxis;
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XXEXEX

23-MAY-2001 AAF94818;

(first entry)

AAF94818 standard; cDNA; 396

ΒP

Human ovarian cancer associated coding sequence SEQ ID

NO: φ

ovarian cancer; vaccine; gene therapy; carcinoma; ss

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AAF94818
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities. Included in the invention polypnucleotide sequences AAF64167 - AAF64175 and peptide AAB75505 where used in the isolation, identification and characterisation of the proteins of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischaemia, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1614
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin ageing due to sunburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expressed, examples of these activities include, immunosuppressive; antiarthritic; antirheumatic; antiproliferative; cytostatic; cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The proteins and polynucleotides, their agonists and antagonists have activities dependent on the tissues and cells in which they are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human secreted proteins AAB75506 - AAB75534 are encoded by polynucleotide sequences AAF64176 - AAF64224. The specification includes amino acid sequences AAB75555 - AAB75606 which represent fragments of the human secreted proteins, and protein sequences with which they share homology.
                                                                                                                                                         361
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                                                                                                             CACCATCCT 1307
                                                                                                                                                         CACCATCCT 369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC
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98.6%;
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Pred. No. 2.4
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated polypeptides associated with ovarian carcinomas, nucleic acids that encode them, useful for the prevention treatment of ovarian cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-SEP-1999; 99US-0394374
01-MAY-2000; 2000US-0561778
15-AUG-2000; 2000US-0640173
07-SEP-2000; 2000US-0656668
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                                      ABT03085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 5; Page 119; 189pp; English.
              ABT03085;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       present sequence is an example of one of these sequences
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                                                                                                                          ACCACCATCCT 369
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                396 BP; 90 A; 117 C; 95 G; 92 T; 2 other;
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1.1e-102;
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Matches 359
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03-APR-2001;
02-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                The present invention provides human ovarian cancer associated and coding sequences. The sequences can be used in the diagnosi treatment of ovarian cancers. The present sequence is a coding
                                                                                                                                                                                                                                                                                                                                                                                                                  Novel ovarian cancer polypeptide and polynucleotide, useful detecting the presence of ovarian cancer in a patient, and i pharmaceutical compositions, e.g. vaccines, for treating over
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                                                                                                                                                                                                                                                                                                                                         of the invention
                                                                                                                                                                                                                                                                                                                                                                                        Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human ovarian carcinoma associated coding
371
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                                                                                                                                                                    GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA 180
                                     CGGGCCAAGGCCCAAGAAAAGGGGAA--GTTCTGCCTCGGCCCTCANGCCATGGCTCCGC
                                                                                                                            CCGCAAGTCCTGTGCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT
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                                                                                                                    CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT
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                  ACCACCATCCT
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359; Conserv
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; 2001US-0825294.
; 2001US-0970966.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene;
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381
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96.8%;
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                                                                                                                                                                                                                                                                                        Score 343.4;
Pred. No. 1.1
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                                                                                                                                                                                                                                                                                                                      2 other;
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completed:

November

7,

2002,

10:10:07

Job time : 78.6545 secs

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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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       seq
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1: /cgn2_6/ptodata/1,

2: /cgn2_6/ptodata/1,

3: /cgn2_6/ptodata/1,

4: /cgn2_6/ptodata/1,

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6: /cgn2_6/ptodata/1,
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/pcTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/backfiles1.seq:*
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US-09-443-113-156-37
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8184.096 Million cell updates/sec
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Sequence 2, Appli
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Sequence 37, Appl
Sequence 37, Appl
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RESULT 2
US-08-484-126-6
US-08-484-126-6
; Sequence 6, Application US/08484126
; Patent No. 5985655
; GENERAL INFORMATION:
; APPLICANT: Anderson, W. French
APPLICANT: Baltrucki, Leon F.

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| ! | P 6                                                            | Db 04                                                                       | Db Qy                                                                  | <b>3</b> 8 60                                                    | us-                                                        |                                         |                                                                             |                                                                            |                                                                               |                                                | RES<br>US-                      |            | 0.0                                                           | 0000                                     | 0000                              | 200                              | 3                                                              | ဂ ဂ                                      |                   |
|---|----------------------------------------------------------------|-----------------------------------------------------------------------------|------------------------------------------------------------------------|------------------------------------------------------------------|------------------------------------------------------------|-----------------------------------------|-----------------------------------------------------------------------------|----------------------------------------------------------------------------|-------------------------------------------------------------------------------|------------------------------------------------|---------------------------------|------------|---------------------------------------------------------------|------------------------------------------|-----------------------------------|----------------------------------|----------------------------------------------------------------|------------------------------------------|-------------------|
|   | 323                                                            | 263<br>268877                                                               | 203<br>268883                                                          | Query M<br>Best Lo<br>Matches                                    | OTHER<br>OTHER<br>OTHER<br>OF 103                          | LENGT<br>TYPE:<br>ORGAN                 | CURRI                                                                       | TITLE<br>TITLE<br>FILE                                                     | APPLI<br>APPLI                                                                | Patent<br>GENERA<br>APPLI                      | SULT 1<br>-09-103-<br>sequence  |            | 444                                                           | 440                                      | 387                               | 4 10 4                           | ມພພ                                                            | 30                                       | 28                |
|   | 23 GGAAGTTCTGCCT 335<br>         <br>717 ATCAGTGCAACCT 2688705 | 7                                                                           | 7 – 6                                                                  | atch 9.<br>cal Similarity 53.<br>71; Conservative                | INFORMATION: CDC INFORMATION: "n" INFORMATION: repr-840A-2 | H: 4403765<br>DNA<br>ISM: Mycobacterium | CURRENT FILING DATE: 1998 NUMBER OF SEQ ID NOS: 2 SOFTWARE: PatentIn Ver. 2 | OF INVENTION: DNA OF INVENTION: TUI REFERENCE: 24366-20 NT APPLICATION NUM | APPLICANT: WHITE, OWEN R. APPLICANT: FRASER, Claire APPLICANT: VENTER, John C | NO. 6294328 L INFORMATION: CANT: FLEISCHMAN, F | 3-840A-2/c<br>ce 2, Application |            | 7.2 7.7.2 7.2 7.2 7.2 7.2                                     | 7.4 7.4 7.4 7.2 7.4                      | 7.4                               | 7.4 7.4                          | 7.6 7.5<br>7.6 7.5<br>7.4 7.4                                  | 7.6 7.5                                  | 7.8 7.5 8         |
| • | 705                                                            | TCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTAACGGGCCCAAGGCCCAAGAAAAGG               | GCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAAC           | 2%; Score 33.8; DB 4;<br>4%; Pred. No. 7.8;<br>0; Mismatches 62; | 1551<br>bases at various positions<br>esent a, t, c or g   | tuberculosis                            | 06-24                                                                       | TUBER: US/09/103.840A                                                      |                                                                               | ert D.                                         | US/09103840A                    | ALIGNMENTS | 4 US-09-393-554-10<br>4 US-09-041-886-20<br>2 US-08-779-801-5 | 5 PCT-US93-12687-1<br>3 US-09-154-083-25 | 2 US-08-826-885-16<br>6 5455158-2 | US-08-410-540-4 PCT-US95-09819-6 | 4 US-08-991-944-3<br>4 US-09-784-316-1<br>4 US-09-404-879A-313 | 4 US-09-423-233-27<br>4 US-09-423-233-26 | 4 US-08-766-528-1 |
|   |                                                                | CCAAGGCCCAAGAAAAGG 322<br>                  <br> CCGACGAGCACGCCGATG 2688718 | CCCCAGGAAACTGAAC 262<br>               <br> CCCCCCTGGAACCCGAAT 2688778 | Length 4403765; ; Indels 0; Gaps 0;                              | s throughout the sequence                                  |                                         |                                                                             | SIS IN MYCOBACTERIUM                                                       |                                                                               |                                                |                                 |            | Sequence 10, Appl<br>Sequence 20, Appl<br>Sequence 5, Appli   | 52,1                                     | . 16                              | 4,04                             | Sequence 3, Appli<br>Sequence 1, Appli<br>Sequence 313, App    | 27, App<br>26, App                       | Sequence 1, Appli |

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US-08-897-443-2/c
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INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
1 FNGTH: 759 bases
          Sequence 2, Application US/08897443
PATENT NO. 5981263
GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                  Matches
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APPLICATION NUMBER: US/08/484,1
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/326,347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Lillie, Raymond J.778
REGISTRATION NUMBER: 31,778
REFERENCE/DOCKET NUMBER: 27:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Lillie, Raymond T
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                123 CCTATCACCAATGCCACCCTGGACCAGC 150
                                                                                                                                                                                                                                                                                                                220 CTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTG 279
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PAGES: 50,
                                                                                                                                                                               340 CCTCANGCCATGGCTCCGCACCACCATC 367
                                                                                                                                                                                                                                             280 CTGCAACACCCCTCTTTGTAACGGGCCCAAGGCCCCAAGAAAAGGGGGAAGTTCTGCCTCGGC 339
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FILING DATE: 20-0CT-1994
APPLICATION NUMBER: 08/973,307
FILING DATE: 09-NOV-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Carella, Byrne, Bain, Gilfillan
ADDRESSEE: Cecchi, Stewart & Olstein
STREET: 6 Becker Farm Road
                                                                                                                                                                                                                                                                                                                                                                 Local
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RESULT 4
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GENERAL INFORMATION:
APPLICANY: Pan, Yang
TITLE OF INVENTION: NOVEL MOLECULES OF TANGO-77 RELATED PROTEIN FAMILY
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: 09404/052001
CURRENT APPLICATION NUMBER: US/09/128,155
CURRENT FILING DATE: 1998-08-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 97; Conserv
                                                                                                                                   Sequence 16, Application US/09128155 Patent No. 6117654
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 415-845-4166 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: PF
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FastSEQ for CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Kaser, Mathew TITLE OF INVENTION: HUMAN NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                        458 ATGGTGCCCGTGGACAGATGCCGCATCCT 430
                                                                                                                                                                                                                                                                                         341 CTCANGCCATGGCTCCGCACCACCATCCT 369
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      161 CAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCC
                                                                                                                                                                                                                                                                                                                                                                281 TGCAACACCCCTCTTTGTAACGGGCCCAAGGCCCCAAGAAAAGGGGGAAGTTCTGCCTCGGCC 340
                                                                                                                                                                                                                                                                                                                                                                                                                                               221 TCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGC 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LIBRARY: UTRS
CLONE: 681719
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE
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3174 Porter Drive
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ilarity 46.4%;
Conservative
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Best Local Similarity 49.4%;
Matches 79; Conservative
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LENGTH: 152331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Pan, Yang
TITLE OF INVENTION: NOVEL MOLECULES OF TANGO-77 RELATED PROTEIN FAMILY
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: 09404/052001
CURRENT APPLICATION NUMBER: US/09/128,155
                                                                                                                                                                                                                                                                                                                                                                                                                                     EARLIER APPLICATION NUMBER: US 60/091,650
EARLIER FILING DATE: 1998-07-02
EARLIER APPLICATION NUMBER: US 60/054,646
EARLIER FILING DATE: 1997-08-04
NUMBER OF SEQ ID NOS: 18
                                                                                                                                                                                                                                                                      ORGANISM: Homo saplens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(176373)
OTHER INFORMATION: n = A,T,C or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/128,155
CURRENT FILING DATE: 1998-08-03
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EARLIER FILING DATE: 1998-07-02
EARLIER APPLICATION NUMBER: US 60/054,646
EARLIER FILING DATE: 1997-08-04
                                    77454 TTGGGGTCGGAGTACATTTGGGGATGAGGGGGAGCTGTAAAAGCAGACTTGGACCTTTTCT 77513
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SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 176373
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LOCATION: (1)...(152331)
OTHER INFORMATION: n = A,T,C or G
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156 TGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGC 195
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                                                          36 TTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTGCTCCT 95
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Pred. No. 23;
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Pred. No. 22;
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CURRENT FILING DATE: 1999-11-19
NUMBER OF SEQ ID NOS: 138
SOFTWARE: PERL Program
SEQ ID NO 67
LENGTH: 327
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                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Yue, Henry
APPLICANT: Baughn, Mariah
APPLICANT: Azimzai, Yalda
APPLICANT: Lal, preeti
TITLE OF INVENTION: MAMMALIAN TOXICOLOGICAL RESPONSE MARKERS
FILE REFERENCE: PC-0007 US
                                                                                  APPLICANT: Wang, Xiaodor APPLICANT: Goldstein, JC TITLE OF INVENTION: METH TITLE OF INVENTION: TO STITLE OF INVENTION: PROPONUMBER OF SEQUENCES: 64 CORRESSONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Kaser, Matthew R. APPLICANT: Panzer, Scott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Zweiger, Gary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: unsure LOCATION: 133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         77574 AGGAGCAGGAGGAGAAGAAGATGGATCTTAAGTGATTTGC
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                                                                                                                                                                                                           APPLICANT: Brown, Michael S. APPLICANT: Briggs, Michael F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              150 AAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCT 209
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                                                                                                                                                                                                                                                                                                                                                                                                               270
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COUNTRY: U.S.A.
                    STATE:
                                                    STREET:
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                                                                     ADDRESSEE:
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                                 Houston
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                  Texas
                                                  P.O. Box 4433
                                                                                                                GOLDSTEIN, JOSEPH L.
VENTION: METHODS AND COMPOSITIONS RELATING
VENTION: TO STEROL REGULATORY ELEMENT BIND:
VENTION: PROTEINS
                                                                                                                                                                                        Briggs, Michael
Wang, Xiaodong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                     Arnold,
                                                                                                                                                                                                         Michael R.
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Pred. No. 1.4;
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Patent No. 5484732
GENERAL INFORMATION:
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                                                                                                        CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: HOWELLS, Stacy L.
REGISTRATION NUMBER: 34,842
REFERENCE/DOCKET NUMBER: PD-2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 455-5100
                                    TELEFAX: (619) 455-5110
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 2205 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "DNA"
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                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 19930322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Rosenfeld, M. G. APPLICANT: Andersen, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1771 AGATGGCCCTGGCTGGCCCAGTGGCTGCCCCC 1805
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                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       262 CTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTAACGGGCCCAAGGCCCAAGAAAAG 321
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LOCATION: 167..3607
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Parker, David L. REGISTRATION NUMBER: 3
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CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC COMPAGE TO COMPAGE PC-DOS/MS-DOS OPERATING SYSTEM: PC-DOS/MS-DOS PC-TO PAPERS #1.0
                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
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STRANDEDNESS:
                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                              90067
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                  NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                              California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Spensley Horn Jubas & Lubitz
1880 Century Park East - Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                              USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TRANSCRIPTION FACTOR FOR THE REGULATION OF THE DEVELOPMENT OF SKIN AND HAIR
                                                                                                                                                                                                                                                                         US/08/035,392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      32,165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 30; DB;
Pred. No. 5.4;
0; Mismatches
                                                                                                                                                           PD-2607
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2; Length 4154;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    41;
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1711 CTCAGATACCACCAGCGTCTACCATAGCCCTGGGCGCAACGTGCTGGGCACCGAGAGCAG 1770
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                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Goldstein, Joseph L.
TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING
TITLE OF INVENTION: TO STEROL REGULATORY ELEMENT BINDING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1771 AGATGGCCCTGGCTGGCCCAGTGGCTGCTGCCCC 1805
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             322 GGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCC 356
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/131,365B
APPLICATION NUMBER: US/08/668,123 FILING DATE: 14-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                   White & Durkee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 30; DB 1
Pred. No. 5.4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 4154;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  41;
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FILING DATE: 01-OCT-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08/131,365
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밁 Ş В US-08-131-365B-37

NAME/KEY: CDS LOCATION: 167..3607

FEATURE:

MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "DNA"

0

TOPOLOGY: STRANDEDNESS:

linear

single

TELEFAX: (512) 474-7577
INFORMATION FOR SEQ ID NO: 37:
SEQUENCE CHARACTERISTICS:
LENGTH: 4154 base pairs

nucleic acid

TELECOMMUNICATION INFORMATION: TELEPHONE: (512) 418-3000

ATTORNEY/AGENT INFORMATION:

FILING DATE: 01-OCT-1993 CLASSIFICATION: 435

TORNEY/AGENTAL L.
NAME: Parker, David L.
22,165

REGISTRATION NUMBER: 32 REFERENCE/DOCKET NUMBER:

Matches Query Match

Local Similarity

8.1%;

Conservative

US-08-668-123-37

Patent No.

APPLICANT:

J. 589100. , INFORMATION: , CANT: Brown, Michael S. , Triggs, Michael R.

APPLICANT:

Wang, Xiaodong

NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:

COUNTRY:

U.S.A.

STREET:

P.O. Box 4433

Arnold,

Houston Texas

ADDRESSEE:

TITLE OF INVENTION:

PROTEINS

PRIOR APPLICATION DATA:

CLASSIFICATION:

CURRENT APPLICATION DATA:

Sequence 37, Application US/08668123 Patent No. 5891631

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS

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                                                               US-08-504-511A-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 3, Application US/08504511A Patent No. 5561224
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              Query Match
Best Local Similarity
   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                           TELEPHONE: (619) 678-50
TELEFAX: (619) 678-5099
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
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CLONE: Skn-la
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                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: HOWELLS, STACY L.
REGISTRATION NUMBER: 34,842
REFERENCE/DOCKET NUMBER: 07:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 678-5070
                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: """
STREET: """
CITY: La Jolla
CTATE: California
""SA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Rosenfeld, M. G. APPLICANT: Andersen, B.
                                                                                                         FEATURE:
                                                                                                                      CLONE: Skn-la
                                                                                                                                                 MOLECULE TYPE: CDNA
                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                245 TCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAACGGG 304
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                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/O: FILING DATE: 20-JUL-1995 CLASSIFICATION: 435
                                                                             NAME/KEY:
                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Fish & Richardson P.C. STREET: 4225 Executive Square, Suite 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OCATION:
                                                                                                                                                                                                                LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              44;
   44;
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                                                                                                                                                                                             nucleic acid
                                                                                                                                                                                                                2205 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Floppy disk
                                                                                                                                                                               single
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              8.0%;
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Score 29.6; DB Pred. No. 5.4; 0; Mismatches
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                             DB 1;
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   24;
                             Length 2205;
   Indels
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                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (415) 324-0880
TELEFAX: (415) 324-0860
INFORMATION FOR SEQ ID NO: 97:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Gregory Dolganov
TITLE OF INVENTION: Transcri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 46
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA: APPLICATION NUMBER: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                    ORIGINAL SOURCE:
INDIVIDUAL ISOLATE:
                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: CDNA to mRNA HYPOTHETICAL: NO ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Sholtz, Charles K.
REGISTRATION NUMBER: 38,
                                                                                                                                                                                                            1392 GCGTTGGTCTCCTCCTCCAGTTCCCGGCGCTTTTCCTCCACCTTGCGCTTCTCCTCCTGG 1333
1212 AGCTCACTTAGGAACTCCTTCCT 1190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 350
CTTY: Palo Alto
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                                                                                                                                                                                                                                              167 GCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCC 226
                                   347 CCATGGCTCCGCACCACCATCCT 369
                                                                                                     287 ACCCCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGGAAGTTCTGCCTCGGCCCTCANG
                                                                                                                                                            227 GGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTTGCATTTGCATCAGCTGCTGCAAC 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         32 CCAAAGCC 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U
ZIP: 94306
                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
                                                                                                                                        TGGACCCGCTTCAGGTGCTCAAACTTCTCATGGAGCTCCCTTTCCTTCTCCTTCAGCTCC
                                                                    94;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1534 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               double
                                                                                                                                                                                                                                                                                                   7.9%;
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                                                                                                                                                                                                                                                                                                 Score 29.2; D
Pred. No. 6.1;
                                                                                                                                                                                                                                                                                   Mismatches 109;
                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                    Length 1534;
                                                                                                                                                                                                                                                                                   Indels
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RESULT 13
US-08-252-073A-5
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US-08-292-945-5
Sequence 5, Application US/08252073A Patent No. 5767073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 5, Application Patent No. 5585478
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (404) 815-6555
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                             ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL: 1
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (404) 815-6508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/990,
FILING DATE: December 10, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,284
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
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                                                                                               465 TATGGGCCCCGACCAGAGGAGTACGAATTCCTCACTCCAG 504
                                                                                                                               188 TCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCG 227
                                                                                                                                                                                              128 GTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTACCGCAAG 187
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CITY: Atlanta
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH:
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                                                                                                                                                                GTTCAACACACATACCGGACTGGCATGAGAGTGGATAAAGCCACATTCATGGTTGGCAGC 464
                                                                                                                                                                                                                                                                                                          78;
                                                                                                                                                                                                                                                                                                       Similarity 48.878; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid
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1100 Peachtree Street, Suite 2800
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Adra, Chaker
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48.8%;
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                                                                                                                                                                                                                                                                                                       Score 28.8; DB Pred. No. 7.1; 0; Mismatches
                                                                                                                                                                                                                                                                                                        0;
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                                                                                                                                                                        Sequence 5, Application PC/TUS9312074 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                           TITLE OF INVENTION: D4 Gene and Methods of Use Thereof NUMBER OF SEQUENCES: 8
COMPUTER READABLE FORM:
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LENGTH: 1107 base pairs
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APPLICATION NUMBER: US 07/990,337
FILLING DATE: 10-DEC-1992
ATTORNEY/AGENT INFORMATION:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/252,07:
FILING DATE: 1-UN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                       465 TATGGGCCCCGACCAGAGGAGTACGAATTCCTCACTCCAG
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                                                                             MEDIUM TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic Center,
STREET: 1201 West Peachtree Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM:
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(404)873-8795
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SYSTEM: PC-DOS/MS-DOS
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48.8%;
                 Release #1.0, Version #1.25 (EPO)
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D4 Gene and Methods of Use Thereof
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APPLICATION NUMBER: PCT/US93/12074
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/990,337
FILING DATE: 10-DEC-1992
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1107 base pairs
TYPE: nucleic acid
crpannerners: 4:forie

PCT/US93/12074

MOLECULE TYPE: CI HYPOTHETICAL: NO

CDNA single

STRANDEDNESS: si TOPOLOGY: linear

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Db 2200625 AACAC 2200629
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US-09-103-840A-2
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ORIGINAL SOURCE:
ORGANISM: Muri
PCT-US93-12074-5
                                                                                                                                                                                                                                                                                                                                            APPLICANT: FLEISCHMAN, Robert D.
APPLICANT: WHITE, Owen R.
APPLICANT: WHITE, Owen R.
APPLICANT: FRASER, Claire M.
APPLICANT: VENTER, John C.
TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM TITLE OF INVENTION: TUBERCULOSIS
FILE REFERENCE: 24366-20007.00
CURRENT APPLICATION NUMBER: US/09/103,840A
CURRENT FILING DATE: 1998-06-24
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
LENGTH: 4403765
                                                                                                                                                    Query Match
Best Local Similarity
Matches 42; Conserv
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Best Local Similarity 48.8%;
Matches 78; Conservative
                                                                                                                                                                                                                                  ORGANISM: Mycobacterium tuberculosis
FEATURE:
OTHER INFORMATION: CDC 1551
OTHER INFORMATION: "n" bases at various positions throughout the sequence
OTHER INFORMATION: represent a, t, c or g
                                          359 ACCAC 363
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Pred. No. 68;
0; Mismatches
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Search completed: November 7, 2002, 15:58:03 Job time: 3169.83 secs

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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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      11111987654321
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369
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Gapop 10.0 , Gapext 1.0
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      /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*
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/cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*
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| 1 G G G G G G G G G G G G G G G G G G G                                                                                                                                                                                        | RESULT 1  JS-09-825-294-199 Sequence 199, Application US/098252 Patent No. US20020004491A1 GENERAL INFORMATION: APPLICANT: Xu, Jiangchun APPLICANT: Stolk, John A. APPLICANT: Algate, Paul A. APPLICANT: Fling, Steven P. TITLE OF INVENTION: COMPOSITIONS APTITLE OF INVENTION: THERAPY AND D. FILE REFERENCE: 210121.484C5 CURRENT APPLICATION NUMBER: US/09/CURRENT FILING DATE: 2001-04-03 NUMBER OF SEQ ID NOS: 215 CURRENT BEST DATE: 2001-04-03 NUMBER OF SEQ ID NOS: 215 SEQ ID NO 199 TYPE: DNA ORGANISM: Homo sapiens FEATURE: NAME/KEY: misc_feature LOCATION: (1)(369) OTHER INFORMATION: n = A,T,C or G S-09-825-294-199                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |                                                                                                                                                 | 22222222222222222222222222222222222222                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |
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APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Filng, Steven P.
APPLICANT: Filng, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
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Local Similarity 99.2%;
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Pred. No. 1.1e-110;
0; Mismatches 3;
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RESULT 3

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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CJ
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 211
LENGTH: 1619
TYPE: DNA
ORGANISM: Homo sapiens
US-09-825-294-211
APPLICANT: X. Jiangchun
APPLICANT: X. Jiangchun
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CFILE REFERENCE: 21012.1484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOSTMARE: FastSEQ for Windows Version 3.0
SEQ ID NO 214
LENGTH: 1897
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APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Filng, Steven P.
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US-09-825-294-214
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Best Local Similarity
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99.2%;
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                                                                                                                                            OF OVARIAN CANCER
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GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun

APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

APPLICANT: Algate, Paul A.

APPLICANT: Eling, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE

TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 9

LENGTH: 396
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: LCCATION: (1)...(396)
: OTHER INFORMATION: n = A,T,C
US-09-825-294-9
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US-09-825-294-9
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US-09-825-294-214
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                                                           Matches 359;
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LOCATION: (1)...(1897)
OTHER INFORMATION: n = A,T,C
                                                                                                                                                                                TYPE: DNA
ORGANISM: Homo sapien
FEATURE:
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Pred. No. 9.7e-104;
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Pred. No. 1.2e-110;
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SOFTWARE: PastSEQ for Windows Version 3.0 SEQ ID NO 212

LENGTH: 1010

TYPE: Days
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Filing, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN
FILE REPERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 212, Application US/09825294 Patent No. US20020004491A1
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Best Local Similarity
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                                           TCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTA 299
                                                                                                        ACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCT
                                                                                                                                                                                                                   AGTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCA
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                                                                                           ACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCT
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98.4%;
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Pred. No. 5.2e-103;
0; Mismatches 4;
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                                    ; ORGANISM: Homo sapien 
US-09-867-701-4251
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US-09-867-701-4251
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US-09-825-294-213
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APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AN
TITLE OF INVENTION: THERRAPY AND DI
FILE REFERENCE: 210121.484C5
                                                                                                                                             GENERAL INFORMATION:

APPLICANT: Aglate, Paul A.

APPLICANT: Jones, Robert

APPLICANT: Harlocker, Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.497

CURRENT APPLICATION NUMBER: US/09/867,701

CURRENT FILING DATE: 2001-05-29
                                                                                   SEQ ID NO 4251
LENGTH: 430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 213
LENGTH: 480
    Query Match
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                                                                                                                 NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2001-04-03
                                                                TYPE: DNA
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les 198; Conservative
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10912
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215
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Pred. No. 2.3e-55;
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   10;
Length 430;
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APPLICANT: Warren, Wesley C.
APPLICANT: Tao, Nengbing
APPLICANT: Byatt, John C.
APPLICANT: Byatt, John C.
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION AND
FILE REFERENCE: 16511.006/37-21(10288)C
CURRENT FILLING DATE: 2001-09-24
CURRENT FILLING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 15112
SEQ ID NO 3015
LENGTH: 407
TYPE: DNA
ORGANISM: Bos taurus
OTHER INFORMATION: Clone ID: 13-LIB34-071-Q1-E1-D1
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APPLICANT: Warren, Wesley C.
APPLICANT: Tao, Nengbing
APPLICANT: Tao, Nengbing
APPLICANT: Byatt, John C.
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILE REPERENCE: 16511.06/37-21(10298)C
CURRENT APPLICATION NUMBER: US/09/960,352
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 15112
SEQ ID NO 13056
LENGTH: 412
                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                 sequence 13056, Application US/09960352
Patent No. US20020137139A1
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Pred. No. 0.3;
0; Mismatches
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                                                                                                                                                                        RESULT 12
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                                                                                       GENERAL INFORMATION:
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APPLICANT: Mathialagan, Nagappan
APPLICANT: Mathialagan, Nagappan
TITLE OF INVENTION: NUCLEIC ACID AND OTHER MOLECULES ASSOCIATED WITH LACTATION .
TITLE OF INVENTION: MUSCLE AND FAT DEPOSITION
FILE REFERENCE: 15511.006/37-21(10298)C
CURRENT APPLICATION NUMBER: US/09/960,352
CURRENT FILING DATE: 2001-09-24
NUMBER OF SEQ ID NOS: 15112
SEQ ID NO 9456
LENGTH: 423
                                                                                                         Sequence 1886, Application US/09960352 Patent No. US20020137139A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9456, Application US/09960352 Patent No. US20020137139A1 GENERAL INFORMATION:
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Best Local Similarity
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                                                                      APPLICANT: Warren, Wesley C.
                APPLICANT:
                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Warren, Wesley C. APPLICANT: Tao, Nengbing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Bos taurus
OTHER INFORMATION: Clone ID: 41-LIB188-009-Q1-E1-C10
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larren, ....
Tao, Nengbing
Byatt, John C.
Byatt, Jayan, Nagappan
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Pred. No. 0.3;
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Pred. No. 0.3;
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; LOCATION: (1)...(544)
; OTHER INFORMATION: n =
US-09-834-975-262
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; ORGANISM: Bos taurus
; OTHER INFORMATION: Clone ID: 09-LIB188-017-Q1-E1-C9
US-09-960-352-1886
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Bolt, Andrew
APPLICANT: Van Huffel, Christophe
APPLICANT: Van Huffel, Christophe
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE IDENTIFICATION, ASSESSMENT, F
TITLE OF INVENTION: OF HUMAN CANCERS
FILE REFERENCE: MRI-016B
CURRENT APPLICATION NUMBER: US/09/834,975
CURRENT APPLICATION NUMBER: 60/197,538
PRIOR APPLICATION NUMBER: 60/197,538
PRIOR APPLICATION NUMBER: 60/197,538
PRIOR FILING DATE: 2000-04-14
PRIOR FILING DATE: 2000-04-14
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Best Local Similarity 51.4
74; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Lillie, James
                                                                                                                                                                                                                                 Matches
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LENGTH: 458
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                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 544
235 GTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCT
                                             144 TGCATACAATTGGTTTTNNGAATNANAACTGGTCAATTTTTCTGCCTATGNGGGCTACTT
                                                                                                                                                                                 115 TTGCACGGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGAT 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             302
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                                                                                        CATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCA 234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACGACTGCTCCTCCCCGAGTTCATTGTGAATTGCACGGTGAACGTTCAAGACATGTGTC 143
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                                                                                                                                      TTGCATGGTNCCTTTTCAACTCATGGGCCACAGGNGACTCTTANAGATTTATGCCAAAAT 143
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45.0%;
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                                                                                                                                                                                                                                                     Score 31.6; DB Pred. No. 0.73;
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RESULT 15
US-10-033-109-13
US-10-033-109-13
Sequence 13, Application US/10033109
Patent No. US20020142390A1
GENERAL INFORMATION:
APPLICANT: Allen, Stephen M.
APPLICANT: Rafalski, J. Antoni
APPLICANT: Sakal, Hajime
TITLE OF INVENTION: Nitrogen Transport Metabolism
FILE REFERENCE: BB-1210
CURRENT APPLICATION NUMBER: US/10/033,109
CURRENT APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/384,625
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/384,625
PRIOR FILING DATE: EARLIER FILING DATE: 1999-08-27
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; ORGANISM: Arabidopsis thaliana
US-09-770-445-893
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US-09-770-445-893
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SOFTWARE: FastSEQ for
SEQ ID NO 893
LENGTH: 775
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APPLICANT:
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                                                                                                                                                                                                                                                                                                               409 GGAGTGGAGAAGGGAACGGCGGAGCTAAGGCCCAACCAAGTAC 450
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Local Similarity 56.9%;
es 58; Conservative
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Hamilton, Carol M.
Price, Jennifer L.
Raines, Tracy M.
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Rameaka, Joshua G.
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Hurban, Patrick
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Davis, Keith R.
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Haas, William David
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; TYPE: DNA
; ORGANIEM: Triticum aestivum
US-10-033-109-13
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                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/098,248
PRIOR FILING DATE: EARLIER FILING DATE: 28 August 1998
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Microsoft Office 97
SEQ ID. NO 13
                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                           Query Match
                                                                               266 GTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAACGGGCCCAAGGCCCCAAGAA 318
                                                                                                                                       645 GGCTTCCTCTTCCACTGGGGCGTCATCGACTACTGCGGCGGCTACGTCATCCACATCCCC 704
                                                        705 GCCGGCGTCGCCGGCTTCACCGCCGCGTACTGGGTCGGGCCAAGGACCAAGAA 757
                                                                                                                                                                               206 GCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCA 265
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7,
  2002, 17:59:58
                                                                                                                                                                                                                     Score 31.4; DB 12; pred. No. 1.5; 0; Mismatches 51;
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Search completed: November Job time: 19.0364 secs

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Database
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Maximum Match 100%
Listing first 45 s
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Maximum DB seq length: 2000000000
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                                                        IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
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369
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(cgn2_6/ptodata/1/pna/US00_SC_COMB.seq:*
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Copyright (c) 1993 - 2002 Compugen Ltd.
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/cgn2_6/ptodata/1/pna/US07_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US102A_COMB.seg:*
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80: /cgn2_6/ptodata/1/pna
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.

Score

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Description

| 21                | 20               | 19               | 18               | 17               | 16               | 15               | 14               | 13               | 12               | 11               | 10               | ,                 | 00                | 7                | 0                 | · U               | 4-1               | . (               | ) N               | , <sub>1</sub> _   |
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| 365.4             | 365.4            | 365.4            | 365.4            | 365.4            | 365.4            | 365.4            | 365.4            | 365.4            | 365.4            | 365.4            | 365.4            | 365.4             | 365.4             | 365.4            | 367               | 367               | 367               | 367               | 367               | 367                |
| 99.0              | 99.0             | 99.0             | 99.0             | 99.0             | 99.0             | 99.0             | 99.0             | 99.0             | 99.0             | 99.0             | 99.0             | 99.0              | 99.0              | 99.0             | 99.5              | 99.5              | 99.5              | 99.5              | 99.5              | 99.5               |
| 1524              | 1524             | 1524             | 1524             | 1524             | 1524             | 1524             | 1524             | 1524             | 1524             | 1524             | 1524             | 1524              | 1524              | 1524             | 369               | 369               | 369               | 369               | 369               | 369                |
| 42                | 42               | 42               | 42               | 42               | 42               | 42               | 42               | 42               | 42               | 42               | 42               | 42                | 42                | 40               | 42                | 36                | 31                | 28                | 25                | Ļ                  |
| US-10-218-849-15  | US-10-218-784-15 | US-10-218-765-15 | US-10-218-631-15 | US-10-218-612-15 | US-10-216-168-15 | US-10-216-167-15 | US-10-216-166-15 | US-10-216-165-15 | US-10-216-164-15 | US-10-216-163-15 | US-10-216-162-15 | US-10-216-160-15  | US-10-216-159A-15 | US-10-119-480-15 | US-10-212-677-199 | US-09-970-966-199 | US-09-825-294-199 | US-09-713-550-199 | US-09-656-668-199 | PCT-US01-45395-199 |
| Sequence 15, Appl | 15.              | 15.              | 15               | 15,              | 15,              | 15,              | 15               | 15,              | 15,              | 15,              | •                | Sequence 15, Appl | 15,               | 15,              | 199               | Sequence 199, App | Sequence 199, App | Sequence 199, App | 9                 | Sequence 199, App  |

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APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
APPLICANT: Filing, Steven P.
APPLICANT: Molesh, David Alan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THE
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.48401PC
CURRENT APPLICATION NUMBER: PCT/US01/45395
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
LENGTH: 369
                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1
PCT-US01-45395-199
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                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 199, Application PC/TUS0145395 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Corixa Corporation APPLICANT: Xu, Jiangchun
                                                                                                                                                                                             NAME/KEY: misc_feature
LOCATION: 29, 345
OTHER INFORMATION: n =
                                                                                                                                                                                                                             ORGANISM: Homo sapiens FEATURE:
181 CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT
                                                                                                   GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC
                                                            Similarity
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US-10-219-061-15
US-10-219-063-15
US-10-219-065-15
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US-10-219-070-15
US-10-219-073-15
US-10-219-073-15
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LENGTH: 369

TYPE: DNA

OGGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(369)
OTHER INFORMATION: n = A,T,C of US-09-656-668-199
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CURRENT APPLICATION NUMBER: US/09/656,668
CURRENT FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 199
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                        ol Similarity
369; Conserv
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Sequence 199, Application US/09825294

GENERAL INFORMATION:
APPLICANT: XU, Jiangchun
APPLICANT: Xtolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
APPLICANT: Filing, Steven P.
TITLE OF INVENTION: COMPOSTIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA
FEILE REFERENCE: 210121.484C5
CURRENT APPLICATION UNUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 199
LENGTH: 369
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; Sequence 199, Application US/09713550
; GENERAL INFORMATION:
; APPLICANT: Xu, Jiangchun
; APPLICANT: Xtolk, John A.
; APPLICANT: Stolk, John A.
; TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER
; TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER
; FILE REFERENCE: 210121.484C4
; FILE REFERENCE: 210121.484C4
; CURRENT APPLICATION UNMBER: US/09/713,550
; CURRENT FILING DATE: 2000-11-14
; NUMBER OF SEQ ID NOS: 205
; NUMBER OF SEQ ID NOS: 205
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LENGTH: 369
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LOCATION: (1)...(369)
OTHER INFORMATION: n = A,T,C or
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ORGANISM: Homo sapiens
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TYPE: DNA
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                                                                                                                                                 OVARIAN CANCER
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APPLICANT: Fling, Steven F.
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
TITLE REFERENCE: 210.121.484C6
CURRENT APPLICATION NUMBER: US/09/970,966
CURRENT ETLING DATE: 2001-10-02
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
SEQ ID NO 199
LENGTH: 369
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                                                                                                       ; NAME/KEY: misc_feature
, LOCATION: 29, 345
; OTHER INFORMATION: n = A,T,C or G
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Best Local Similarity
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                                                    Matches 369;
                                                                               Query Match
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 Molesh, David Al Fling, Steven P.
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FEATURE:
NAME/KEY: misc_feature
LOCATION: 29, 345
OTHER INFORMATION: n = A,T,C or G
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CURRENT FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 288
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 199
LENGTH: 369
                                                                                                                                                                                                                                                                      Matches
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Chenault, Ruth A.
APPLICANT: Xu, Jiangchun
APPLICANT: Fanger, Gary R.
APPLICANT: Harlocker, Susan L.
APPLICANT: McNeill, Patricia D
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US-10-119-480-15
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LENGTH: 1524
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501 CACCATCCT
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                                                                                                                                                               CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT
                                                                                                                                                                                                                      CCGCAAGTCCTGTGCATCAGCGGCCTGTCTCATCGCCTGTGCGGGTACCAGTCCTT 240
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Godowski, Paul J.
Gorimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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Pred. No. 3.3e-98;

0; Mismatches (
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US-10-216-159A-15

RESULT 8

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FILLE OF INVESTION. OCCUPANTS.
FILL REFERENCE: P3559DIC6

CURRENT APPLICATION NUMBER: US/10/216,159A

CURRENT FILING DATE: 2002-08-09

PRIOR APPLICATION NUMBER: 10/119,480

PRIOR FILING DATE: 2002-04-09

PRIOR APPLICATION NUMBER: 60/059113

PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/062287

PRIOR APPLICATION NUMBER: 60/06287

PRIOR FILING DATE: 1997-10-28

PRIOR FILING DATE: 1997-10-31

PRIOR APPLICATION NUMBER: 60/064103

PRIOR FILING DATE: 1997-10-31

PRIOR APPLICATION NUMBER: 60/064103

PRIOR FILING DATE: 1997-10-31

PRIOR APPLICATION NUMBER: 60/064973

PRIOR APPLICATION NUMBER: 60/069873
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PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/079656
PRIOR FILING DATE: 1998-03-27
PRIOR FILING DATE: 1998-03-27
PRIOR FILING DATE: 1998-03-27
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APPLICANT:
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NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15.
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APPLICANT: Desnoyers, Luc
APPLICANT: Gerritsen, Mai
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ORGANISM: Homo Sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH: 1524
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                                                                                                                                                                                                                  GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA 180
                         CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCAC 360
CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA 300
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Gerritsen, Mary
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Grimaldi, J. Christopher
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       99.0%; Score 365.4; DB 42; 99.2%; Pred. No. 3.3e-98;
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US-10-216-160-15
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NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
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APPLICANT:
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CURRENT APPLICATION NUMBER: US/10/216,160
CURRENT FILING DATE: 2002-08-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
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                                                               301 CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCATGGCTCCGCAC 360
                                                                                                        381
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501 CACCATCCT 509
                        361 CACCATCCT 369
                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                           CCGCAAGTCCTGTGCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT 240
                                                                                                      CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA 300
                                                 Watanabe, Colin L. Wood, William I.
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Smith, Victoria
Stephan, Jean-Philippe F.
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Gerritsen, Mary
Goddard, Audrey
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Grimaldi, J. Christopher
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                                                                                                                                                                                                                                                                                                                                                                                 99.0%; Score 365.4; DB 42; Length 1524; 99.2%; Pred. No. 3.3e-98; tive 0; Mismatches 3; Indels 0;
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Sequence 15, Application US/10216162; GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Desnoyers, Luc

US-10-216-162-15

APPLICANT:

Gerritsen, Mary

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CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/06287
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
PRIOR PRIOR PRIOR DATE: 1997-12-17
PRIOR PRIOR PRIOR DATE: 1997-12-17
PRIOR PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR PRIOR APPLICATION NUMBER: 60/079294
PRIOR PRIOR PRIOR DATE: 1998-03-25
PRIOR FILING DATE: 1998-03-25
PRIOR PRILING DATE: 1998-03-26
PRIOR FILING DATE: 1998-03-26
PRIOR FILING DATE: 1998-03-27
PRIOR FILING DATE: 1998-03-27
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SEQ ID NO 15
LENGTH: 1524
TYPE: DNA
ORGANISM: Homo Sapien
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501 CACCATCCT 509
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                                          361 CACCATCCT 369
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       201
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                                                                      CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCAGGCCCAGGGCTCCGCAC
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                                                                                                                                                         CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA 440
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Grimaldi, J. Christopher
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Smith, Victoria
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US-10-216-163-15
; Sequence 15, Application US/10216163
; GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                         Matches 366;
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NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
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PRIOR FILLYG DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILLYG DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILLYG DATE: 1997-10-17
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PRIOR APPLICATION NUMBER: 60/079656
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PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3530P1C3
CURRENT APPLICATION NUMBER: US/10/216,163
CURRENT FILING DATE: 2002-08-09
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APPLICANT: Desnoyers, Luc
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381 CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA 440
                 241 CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCTGCAACACCCCTCTTTGTAA 300
                                                                                        321
                                                                                                                           181
                                                                                                                                               141 GGCAACTTTTGCGGATTGTTCTTGCTTCCAGGCTTTGCGCTGCAAATCCAGTGCTACCA 200
                                                                                                                                                                                                                                            201
                                                                                                                                                                                                                                                             61 GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC 120
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                                                                      CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT 380
                                                                                            CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT 240
                                                                                                                                                                                                                          Stephan, Jean-Philippe F. Watanabe, Colin L. Wood, William I.
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Smith, Victoria
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Grimaldi, J. Christopher
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Gerritsen, Mary
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                                                                                                                                                                                                                                                                                                                                                                                                 99.0%; Score 365.4; DB 42; 99.2%; Pred. No. 3.3e-98;
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 Mismatches

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CURRENT APPLICATION NUMBER: US/10/216,164
CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-28
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                                                                                                                                                                                                                            Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 15
LENGTH: 1524
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PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/079656
PRIOR APPLICATION NUMBER: 60/079656
                                                                                                                                                                                                                                                                                                                                                                                            Remaining Prior Application data removed - See File Wrapper or PALM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Baker, Kevin APPLICANT: Desnoyers, I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
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                               121 GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA 180
                                                                                                                                            261
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                                                                                                                  61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  t APPLICATION NUMBER: 60/069873
FILING DATE: 1997-12-17
PAPPLICATION NUMBER: 60/078910
FILING DATE: 1998-03-20
                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 1998-03-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 60/079728
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                                                                   GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC 120
GGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGTA
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Grimaldi, J. Christopher
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                                                                                                                                                                                                                                            99.0%;
99.2%;
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                                                                                                                                                                                                                          Score 365.4; DB 42; Length 1524; Pred. No. 3.3e-98; 0; Mismatches 3; Indels 0;
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CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
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                                                                     Matches 366;
                                                                                     Best Local Similarity
                                                                                                     Query Match
                                                                                                                                                                                                                                                         PRIOR FILING DATE: 1998-03-26
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1998-03-27
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PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-31
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                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 1998-03-20 PRIOR APPLICATION NUMBER: 60/
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                                                                                                                                                   TYPE: DNA
ORGANISM: Homo Sapien
     141
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                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 60/079728
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FILING DATE: 1998-03-25
                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/078910
GGCAACTTTTTGCGGATTGTTCTTGCTTCCAGGCTTTGCGCTGCAAATCCAGTGCTACCA 200
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Grimaldi, J. Christopher
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Gerritsen, Mary
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                                                                     Conservative
                                                                                 99.0%;
                                                                   0;
                                                                                 Score 365.4; DB (
                                                                   Mismatches
                                                                                                                                                                                                                                     See File Wrapper or PALM.
                                                                                                   DB 42;
                                                                 Indels
                                                                                                 Length
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; TYPE: DNA
; ORGANISM: Homo
US-10-216-166-15
                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/216,166
CURRENT ETLING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/06287
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/063549
PRIOR APPLICATION NUMBER: 60/064103
PRIOR FILING DATE: 1997-10-28
PRIOR PRIOR FILING DATE: 1997-10-31
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/069873
PRIOR APPLICATION NUMBER: 60/078910
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/079294
PRIOR FILING DATE: 1998-03-25
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GENERAL INFORMATION:
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APPLICANT: Desnoy
APPLICANT: Gerrit
APPLICANT: Goddar
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                                                                                        NUMBER OF SEQ ID
                                                                                                                              PRIOR FILING DATE: 1998-03-27
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                                                                     NO 15
                                                                                                                                                                   APPLICATION NUMBER: 60/079656 FILING DATE: 1998-03-26
                                                                                                                                               APPLICATION NUMBER: 60/079728
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Grimaldi, J. Christopher
Gurney, Austin L.
Smith, Victoria
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Gerritsen, Mary
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                                                                                          Application data removed - D NOS: 246
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                                                                                                           See File Wrapper or
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                                             CURRENT FILING DATE: 2002-08-09
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR FILING DATE: 2002-04-09
PRIOR FILING DATE: 2002-04-09
PRIOR PPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-10-17
PRIOR PPLICATION NUMBER: 60/06287
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-31
PRIOR APPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/069873
PRIOR FILING DATE: 1997-12-17
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15, Application US/10216167 GENERAL INFORMATION:
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Best Local Similarity
Matches 366; Conserv
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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CURRENT APPLICATION NUMBER: US/10/216,167
CURRENT FILING DATE: 2002-08-09
                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Stephan, Jean-Philippe F.
APPLICANT: Watanabe, Colin L.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Baker, Kevin
APPLICANT: Desnoyers, I
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Smith, Victoria
Stephan, Jean-Philippe F.
Watanabe, Colin L.
Wood, William I.
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Gerritsen, Mary
Goddard, Audrey
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                      1998-03-26
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60/079728
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; ORGANISM: Homo Sapien
US-10-216-167-15
Search completed: November 7, 2002, 17:54:11 Job time: 830.855 secs
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Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
                                                                                                                                                                                                                                                                                                                                                   Query Match 99.0%; Score 365.4; DB 42; Length 1524; Best Local Similarity 99.2%; Pred. No. 3.3e-98; Matches 366; Conservative 0; Mismatches 3; Indels 0; Gaps
                                         361 CACCATCCT 369
|||||||||
501 CACCATCCT 509
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Result
No.
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Maximum DB
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score greater than or equal to the score of the result being
and is derived by analysis of the total score distribution.
 Score
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length: 2000000000
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1: /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*

3: /cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*

4: /cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*

5: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*

6: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*

7: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
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369
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                                                                                                                                                                                                                                            Query
Match Length
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Copyright (c) 1993 - 2002
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  first 45 summaries
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   US-10-127-829A-293

US-10-127-831A-293

US-10-127-837A-293

US-10-127-842A-293

US-10-127-842A-293

US-10-127-901A-293

US-10-128-689A-293

US-10-131-833A-293

US-10-131-833A-293

US-10-131-833A-293

US-10-131-833A-293
                                                                                                                       US-10-266-829-18

US-10-281-024-16

US-10-131-813A-293

US-10-131-813A-293

US-10-131-823A-293

US-10-131-824A-293

US-10-131-824A-293

US-10-131-826A-293

US-10-131-829A-293

US-10-125-926A-293

US-10-125-926A-293
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US-10-085-198-105
US-10-085-198-107
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 -10-125-930A-293
-10-127-825A-293
-10-127-838B-293
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| 45                                       | 3 4               | 5 12              | 40                                     | 38                | 37                 | 36                 | 5                  | 34                 | ü                  | 22                 | 3                  | 30                 | 29                 | 8                  | 27                 |
| 27.6<br>27.4                             | 27.6              | ) N<br>B B        | 28<br>28                               | 28.2              | 28.8               | 28.8               | 28.8               | 28.8               | 28.8               | 28.8               | 28.8               | 28.8               | 28.8               | 28.8               | 28.8               |
| 7.5                                      | 7.5               | 7.6               | 7.6                                    | 7.6               | 7.8                | 7.8                | 7.8                | 7.8                | 7.8                | 7.8                | 7.8                | 7.8                | 7.8                | 7.8                | 7.8                |
| 450<br>6227                              | 385               | 6085              | 413<br>6085                            | 470               | 3449               | 3449               | 3449               | 3449               | 3449               | 3449               | 3449               | 3449               | 3449               | 3449               | 3449               |
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| US-09-513-999C-2171<br>US-10-240-425-364 | US-09-620-607B-20 | US-09-484-331-8   | US-09-513-999C-2172<br>US-09-484-743-8 | US-10-240-425-128 | US-10-137-873A-293 | US-10-137-872A-293 | US-10-131-836A-293 | US-10-131-821A-293 | US-10-128-693A-293 | US-10-128-690A-293 | US-10-128-686A-293 | US-10-128-685A-293 | US-10-128-684A-293 | US-10-127-849A-293 | US-10-127-843A-293 |
| Sequence 2171, Ap<br>Sequence 364, App   | 20,               | Sequence 8, Appli | Sequence 21/2, Ap<br>Sequence 8, Appli |                   | Sequence 293, App  |                    | •                  | `                  | -                  | •                  | -                  | -                  | Sequence 293, App  |                    | Sequence 293, App  |

## ALIGNMENTS

RESULT 1

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CURRENT APPLICATION NUMBER: US/10/230,437
CURRENT FILING DATE: 2002-08-28
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR APPLICATION NUMBER: 10/119,480
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/05287
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-10-17
PRIOR PRIOR APPLICATION NUMBER: 60/06287
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-31
PRIOR FILING DATE: 1997-10-31
PRIOR PRIOR APPLICATION NUMBER: 60/069873
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/079294
PRIOR APPLICATION NUMBER: 60/079294
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                                         ; ORGANISM: HOMO US-10-230-437-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Gurney, Austin L.

APPLICANT: Smith, Victoria

APPLICANT: Stephan, Jean-Philippe F.

APPLICANT: Watanabe, Colin L.

APPLICANT: Wood, William I.

TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES

FILE REFERENCE: P3530P1C94

CURRENT APPLICATION NUMBERS OF THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15, Applica GENERAL INFORMATION:
                                                                                                       Remaining Prior Application
NUMBER OF SEQ ID NOS: 246
SEQ ID NO 15
LENGTH: 1524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Baker,
                                                                                                                                                                                                                                         PRIOR FILING DATE: 1998-03-25
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1998-03-26
                                                                                                                                                                                             PRIOR FILING DATE: 1998-03-27
                                                                                     TYPE: DNA
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Gerritsen, Mary
Goddard, Audrey
                                                                Sapien
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99.0%;
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                                                                                                                                                                             data removed -
Score 365.4;
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DB
6;
Length 1524;
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Best Local Similarity

99.2%;

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; ORGANISM: Homo sapiens US-10-085-198-105
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       Best Loc
Matches
                                                                                                                               SEQ ID NO 105
LENGTH: 4321
                                       Query Match
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TITLE OF INVENTION: Proteins and Nucleic Acids
FILE REFERENCE: 21402-279
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PRIOR TILING DATE: 2001-08-13
PRIOR APPLICATION NUMBER: 60/312,858
PRIOR FILING DATE: 2001-08-16
PRIOR APPLICATION NUMBER: 60/271,840
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CURRENT FILING DATE: 2002-02-25
                                                                                                                                                                                    Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 653
                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/272,405 PRIOR FILING DATE: 2001-02-28
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PRIOR APPLICATION NUMBER: 60/315,614
PRIOR FILING DATE: 2001-08-29
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PRIOR APPLICATION NUMBER: 60/299,695
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PRIOR APPLICATION NUMBER: 60/277,324
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PRIOR FILING DATE: 2001-02-26
Local 5.71;
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                    Similarity
                                                                                                                                                                   PatentIn Ver.
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   8.5%;
ilarity 51.4%;
Conservative
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 ; Score 31.4; D; Pred. No. 1.2; O; Mismatches
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PRIOR APPLICATION NUMBER: 60/271,646
PRIOR APPLICATION NUMBER: 60/276,401
PRIOR APPLICATION NUMBER: 60/276,401
PRIOR APPLICATION NUMBER: 60/276,401
PRIOR APPLICATION NUMBER: 60/311,981
PRIOR FILING DATE: 2001-08-13
PRIOR APPLICATION NUMBER: 60/312,858
PRIOR APPLICATION NUMBER: 60/312,858
PRIOR APPLICATION NUMBER: 60/271,840
PRIOR FILING DATE: 2001-08-16
PRIOR APPLICATION NUMBER: 60/271,840
PRIOR FILING DATE: 2001-02-27
PRIOR FILING DATE: 2001-02-27
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Best Local Similarity
Matches 71; Conserv
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SOFTWARE: PatentIn Ver. 2
SEQ ID NO 107
LENGTH: 7097
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PRIOR FILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: 60/286,096
PRIOR FILING DATE: 2001-04-21
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PRIOR FILING DATE: 2001-08-29
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PRIOR FILING DATE: 2001-06-20
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                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapiens
  3676
                                                                                             3616 TCCCGGCTCTCACCTCTGACCCAAAGCCCCCTCAGCTCCCGCACCGGCTCCCCTGAGCTC 3675
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                                              352 GCTCCGCACCACCATCCT 369
                                                                                                                                                                                                                      232 CCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCCAGTTTGCATCAGCTGCTGCAACACCCC 291
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GCCGCCCGTGCCCGGCCT 3693
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RESULT 4 US-10-266-829-18

Sequence 18, Applica GENERAL INFORMATION:

Application US/10266829

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CURRENT APPLICATION NUMBER: US/10/266,829

CURRENT FILING DATE: 2002-10-09

PRIOR APPLICATION NUMBER: 09/756,168

PRIOR FILING DATE: 2001-01-09

PRIOR FILING DATE: 2000-07-20

PRIOR FILING DATE: 2000-07-20

PRIOR APPLICATION NUMBER: 60/145,220

PRIOR APPLICATION NUMBER: 60/145,220

PRIOR FILING DATE: 1999-07-23
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Best Local S
Matches 56
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LENGTH: 1372
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                                                                                                                                      Matches
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APPLICANT: KYOKO HIGUCHI
TITLE OF INVENTION: NICCTIANAMINE SYNTHASE AND GENE ENCODING
TITLE OF INVENTION: THE SAME
FILE REFERENCE: 55107 (71526)
CURRENT APPLICATION NUMBER: US/10/281,024
CURRENT FILING DATE: 2002-10-25
PRIOR APPLICATION NUMBER: US/09/674,337A
PRIOR PILING DATE: 2002-01-07
PRIOR PILING DATE: 2002-01-07
PRIOR PILING DATE: 2002-01-07
PRIOR APPLICATION NUMBER: PCT/JP99/02305
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SOFTWARE: PatentIn Ver. 2
SEQ ID NO 18
                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 199
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TITLE OF INVENTION: 29 Human secreted proteins
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OTHER INFORMATION: n equals a,t,g,
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158 GAGCAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCAGCGGGCC
                                                        835 CCCGAGGACGTCAGGCGTGGCGGGTTCGACGTTCTGGCGGTGTGCCACCCGGAGGACGAG 894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 455 TCTTGGGGGCTTGTTGATTGGTGCTGGCTGTGCCCCTCTACCCCTTGGGCTGGGACAGTGA 514
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                                                                                          Local Similarity
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    Mismatches

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Pred. No. 2.
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US-10-131-813A-293/c
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LENGTH: 3449
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Best Local Similarity
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667 CGGGCCCCCTCTGCTTCTGAGAATGCGATGTTCAGGGCATACTGGATGGCCAGCCCAGTC
                                  281 TGCAACACCCCTCTTTGTAACGGGCCAAGGCCCAAGAAAAGGGGGAAGTTCTGCCTCGGCC
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                                                                                                               TCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGC 280
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45.9%;
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Pred. No. 7.5;
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US-10-131-819A-293/c

Sequence 293, Application US/10131819A

GENERAL

APPLICANT: Baker, Kevin

INFORMATION

APPLICANT: APPLICANT:

DeForge, Laura Beresini, Maureen

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                                                                                                                                       Matches
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Best Local
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CURRENT APPLICATION NUMBER: US/10/131,823A
CURRENT FILING DATE: 2002-04-24
PRIOR ADDITION:
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PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-17
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                                 221 TCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGC 280
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TCCTGAGGTCTCCCATCTGTCACGATCATTATGACCCGTGGCACATTCTCCCTCAGGGGC
                                                                   CCAATGGCAAAGATTAGGATGCCCGTGTCCCGTGCCTTAGCAGCCACCTCGGCCACGGAG 728
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Pred. No. 7.
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    See File Wrapper

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망 QΥ рь Š S US-10-131-819A-293 Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 550
SEQ ID NO 293
LENGTH: 3449 Matches Query Match Best Local Similarity PRIOR APPLICATION NUMBER: 60/056974 PRIOR FILING DATE: 1997-08-26 PRIOR APPLICATION NUMBER: 60/059113 PRIOR APPLICATION NUMBER: 60/0 PRIOR FILING DATE: 1997-06-18 CURRENT APPLICATION NUMBER: US/10/131,819A CURRENT FILING DATE: 2002-04-24 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME APPLICANT: APPLICANT: APPLICANT: APPLICANT: PRIOR APPLICATION NUMBER: 60/059122 PRIOR FILING DATE: 1997-09-17 PRIOR FILING DATE: 1997-09-17 PRIOR APPLICATION NUMBER: 60/ APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: PRIOR APPLICATION NUMBER: 60/059115 PRIOR FILING DATE: 1997-09-17 FILE REFERENCE: P3330R1C134 APPLICANT: APPLICANT: PRIOR FILING DATE: 1997-09-19 PRIOR PRIOR FILING DATE: 1997-09-17 TYPE: DNA ORGANISM: Homo 281 TGCAACACCCCTCTTTGTAACGGGCCAAGGCCCAAGAAAGGGGGAAGTTCTGCCTCCGCC 727 221 TCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGC 787 CCAATGGCAAAGATTAGGATGCCCGTGTCCCGTGCCTTAGCAGCCACCTCGGCCACGGAG 161 CAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCC 220 APPLICATION NUMBER: 60/059352 FILING DATE: 1997-09-19 APPLICATION NUMBER: 60/ FILING DATE: 1997-09-18 FILING DATE: 1997-09-17 APPLICATION NUMBER: 60/059184 APPLICATION NUMBER: TCCTGAGGTCTCCCATCTGTCACGATCATTATGACCCGTGGCACATTCTCCCTCAGGGGC Zhang, Zemin Smith, Victoria Goddard, Audrey Godowski, Paul J. Gurney, Austin L. Gerritsen, Mary E. Gao, Wei-Qiang Desnoyers, Luc Filvaroff, Ellen Wood, William Watanabe, Colin K Stewart, Timothy A. Sherwood, Steven Conservative 7.8%; 45.9%; 60/056974 60/059263 60/049911 60/059588 60/059117 Score 28.8; DB 6; Length 3 Pred. No. 7.5; Indels 0; Mismatches 113; Indels Length 3449; 0; Gaps 340 280 668 728

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Best Local
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LENGTH: 3449
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GENERAL INFORMATION
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CURRENT APPLICATION NUMBER: US/10/131,824A
CURRENT FILING DATE: 2002-04-24
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES
TITLE OF INVENTION: ACIDS ENCODING THE SAME
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PRIOR FILING DATE: 1997-08-26
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221 TCTGCCGGGTACCAGTCCTTCTGCTCCCCAGGGAAACTGAACTCAGTTTTGCATCAGCTGC 280
                                                                            161 CAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCC 220
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                                                       CCAATGGCAAAGATTAGGATGCCCGTGTCCCGTGCCTTAGCAGCCACCTCGGCCACGGAG 728
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Godowski, Paul J.
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Pred. No. 7.
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US-10-131-826A-293/c
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PRIOR FILING DATE: 1997-08-26
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                   161 CAAAGTGCCGGGATCATGTACCGCAAGTCCTGTGCATCAGCGGCCTGTCTCATCGCC 220
787 CCAATGGCAAAGATTAGGATGCCCGTGTCCCGTGCCTTAGCAGCCACCTCGGCCACGGAG 728
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Godowski, Paul J.
Gurney, Austin L.
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Wood,William
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Filvaroff, Ellen
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US-10-131-829A-293
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US-10-131-829A-293/c
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                                                                                             Query Match
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PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
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NUMBER OF SEQ ID NOS: 550
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                                                                           Local
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Godowski, Paul J.
Gurney, Austin L.
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Watanabe, Colin K
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                                                                         ; TYPE: DNA; Homo Sapien US-10-125-926A-293
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US-10-125-926A-293/c
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                                                                                                                                Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 293
LENGTH: 3449
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 Matches
               Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/125,926A CURRENT FILING DATE: 2002-10-15
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PRIOR FILING DATE: 1997-09-17
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PRIOR APPLICATION NUMBER: 60/059117
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                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/059184
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FILING DATE: 1997-09-17
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Conservative
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               7.8%;
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US-10-127-829A-293/C
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  Query Match
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                                                                                                                   SEQ ID NO 293
LENGTH: 3449
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                                                                                                                                                        Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 550
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C85
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                                                                                                   TYPE: DNA
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                                                                                                                                                                                             FILING DATE: 1997-09-19
APPLICATION NUMBER: 60/059588
FILING DATE: 1997-09-19
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APPLICATION NUMBER: 60/059352
                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/059184 FILING DATE: 1997-09-17
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FILING DATE: 1997-09-17
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Sherwood, Steven
Smith, Victoria
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Godowski, Paul J.
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Gerritsen, Mary E.
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PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
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              ORGANISM: Homo Sapien
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                                                     LENGTH:
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                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/059122
FILING DATE: 1997-09-17
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FILING DATE: 1997-09-17
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Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
Smith, Victoria
Stewart, Timothy A.
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Wood, William
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Filvaroff, Ellen
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US-10-127-835A-293

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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C102
CURRENT APPLICATION UMBER: US/10/127,835A
CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-08-18
PRIOR FILING DATE: 1997-08-26
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059117
PRIOR APPLICATION NUMBER: 60/059117
PRIOR APPLICATION NUMBER: 60/059117
PRIOR APPLICATION NUMBER: 60/059117
PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
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US-10-127-835A-293/c
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GENERAL INFORMATION:
APPLICANT: Baker Kevin P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 7.8%;
Best Local Similarity 45.9%;
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                       SEQ ID NO 293
LENGTH: 3449
TYPE: DNA
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PRIOR
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PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
                                                                              Remaining Prior Application data NUMBER OF SEQ ID NOS: 550
                                                                                                                        PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
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ORGANISM: Homo Sapien
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                                                                                                                                                                                                                                                               APPLICATION NUMBER: 60/059184 FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/059122 FILING DATE: 1997-09-17
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Watanabe, Colin K
Wood, William
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Godowski, Paul J.
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                                                               CGGGCCCCCTCTGCTTCTGAGAATGCGATGTTCAGGGCATACTGGATGGCCAGCCCAGTC
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                                                                                                                              TCCTGAGGTCTCCCATCTGTCACGATCATTATGACCCGTGGCACATTCTCCCTCAGGGGC 668
ATGGTGCCCGTGGACAGATGCCGCATCCT 579
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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## REFERENCE AUTHORS TITLE KEYWORDS SOURCE ORGANISM RESULT 1 AC079773/c LOCUS DEFINITION Result No. VERSION ACCESSION a O O ი 000 a Homo sapiens AC079773 AC079773.8 G Eukaryota; Metazoa; Chordata; Craniata; Mammalia; Eutheria; Primates; Catarrhin; 1 (bases 1 to 129676) Sulston, J.E. and Waterston, R. Toward a complete human genome sequence Genome Res. 8 (11), 1097-1108 (1998) Homo sapiens. Homo sapiens Query Match Length 3.3 444 3.3 92874 9.8 970395 9.7 167693 7.7 16967 7.7 180837 7.7 184485 7.7 192464 7 200145 1388 1388 1388 2666 3190 3190 3190 394029 99020 994029 99540 110000 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 111914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 11914 119 GI:15145561 129676 bp DNA linear PRI 09-JA BAC clone RPI1-258B17 from 2, complete sequence. DB 10 AC126575 AC098517\_0 AL596180 AC103498 AC094168 AC123419 ATHAP2A AY065052 AY065057 AB035577 AB019231 AC114102 ATT24H18 AC117309 ATT2L20 AC096152 AC107560 AC102588 AC123116 AC094585 AC022824 AC009679 AC090971 AC126666 AX393645 AC112072 AC099984 AC026770 AC011639 AC022742 AX319944 AX150120 AX362255 AX136698 AX319942 AF034633 ALIGNMENTS SUMMARIES Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo. AX319944 Sequence AX150120 Sequence AX150120 Sequence AX150120 Sequence AX150120 Sequence AX150120 Sequence AC099984 Mus muscu AC029770 Homo sapi AC011639 Homo sapi AC012742 Homo sapi AC0209671 Homo sapi AC0209671 Homo sapi AC0209672 Homo sapi AC0209672 Homo sapi AC0209673 Homo sapi AC126666 Rattus no AX393645 Sequence Y13720 Arabidops AY065052 Arabidops AY065052 Arabidops AY065052 Arabidops AX1924474 Aublyoma AB019231 Arabidops AC103197 Aublyoma AB019231 Arabidops AC117309 Rattus no AL353013 Arabidops AC117309 Rattus no AL596180 Mouse DNA AC103498 Rattus no AC024168 Rattus no AC024168 Rattus no AC021823 Homo sapi AC012588 Mus muscu AC102588 Mus muscu AC103116 Rattus no AC102588 Mus muscu AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC103116 Rattus no AC079773 BC011449 Description PRI 09-JAN-2002 Sequence Sequence Sequence Sequence Sequence Sequence

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Submitted (09-AUG-2001) Genome
University School of Medicine,
MO 63108, USA
5 (bases 1 to 129676)
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Submitted (10-SEP-2000) Genome Sequencing Center, Washington
University School of Medicine, 4444 Forest Park Parkway, St. Louis,
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The sequence of Homo sapiens BAC clone RP11-258B17
Unpublished (2001)
3 (bases 1 to 129676)
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                                                                                                                                                                                                                                                                                                                                                      The RPCI-11 human BAC library was made from the blood of one male donor, as described by Osoegawa, K., Woon, P.Y., Zhao, B., Frengen, E., Tateno, M., Catanese, J.J. and de Jong, P.J. (1998) An improved approach for construction of bacterial artificial chromosome libraries. Genomics 51:1-8. The clone may be obtained either from Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong and coworkers at the Roswell Park Cancer Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.
                                                                                                                                                                           NEIGHBORING SEQUENCE INFORMATION: The clone sequenced to the left is RP11-1L22; the clone sequenced to the right is RP11-159N20, 2000 bp overlap. Actual start of the clone is at base position 1 of RP11-258B17; actual end is at base position 60003 of RP11-159N20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOURCE INFORMATION: The RPCI-11 human B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mapping information for this clone was provided by Dr. John D. McPherson, Department of Genetics, Washington University, St. Louis MO. For additional information about the map position of this sequence, see http://genome.wustl.edu/gsc
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Contact: sapiens@watson.wustl.edu
----- Summary Statistics
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/organism="Homo sapiens"
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15532
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12697. .12739
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1397. .3331
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27048.
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26656.
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25108
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20170. .20305
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                          TGGGCAGACACGAGCAGGAGGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAG
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7. .32787
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                                                                                                                                                                                                                                                                                                                                              Clone distribution: MGC clone distribution information can be fo through the I.M.A.G.E. Consortium/LLML at: http://image.llnl.gov Series: IRAL Plate: 25 Row: n Column: 18
This clone was selected for full length sequencing because it passed the following selection criteria: Hexamer frequency ORF
                                                                                                                                                                                                                                                                                                                                                                                                                                                        CDNA Library Preparation: Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LINL)
DNA Sequencing by: Institute for Systems Biology
http://www.systemsbiology.org
contact: amadan@systemsbiology.org
Anup Madan, Rachel Dickhoff, Jessica Fahey, Stephanie Ford, Julia
Greene, Mark Ketteman and Anuradha Madan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
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Direct Submission
Submitted (25-JUL-2001) National Institutes of Health, Mammalian Submitted (25-JUL-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Gene Collection (MGC), Room 11A03, Bethesda, MD 20892-2590,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 826)
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/tissue_type="5kin, melanotic me
/clone_lib="NIH_MGC_20"
/lab_host="DH10B-R"
/note="Vector: pOTB7"
a 197 c 228 g 164 t
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Direct Submission
Submitted (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team);
Kazusa-Kamateari, Kisarazu, Chiba 292-0812, Japan
(E-mail:genomics@hri.co.jp, Tel:81-438-52-3975, Fax:81-438-52
NEDO human cDNA sequencing project supported by Ministry of
                                                                                                     Kawakami,B., Sugiyama,A., Takemoto,M., Sugiyama,T., Irie,R., Otsuki,T., Sato,H., Wakamatsu,A., Ishii,S., Yamamoto,J., Isono Kawai-Hio,Y., Saito,K., Nishikawa,T., Kimura,K., Yamashita,H., Matsuo,K., Nakamura,Y., Sekine,M., Kikuchi,H., Kanda,K., Wagatsuma,M., Murakawa,K., Kanehori,K., Takahashi-Fujii,A., Oshima,A., Suzuki,Y., Sugano,S., Nagahari,K., Masuho,Y., Nagai and Isogai,T.
                                                                                                                                                                                                                                            clone:BRALZ2001350.
Homo sapiens
                                                                                                                                                                                                                                                                  oligo capping; fis (full insert sequence). Homo sapiens alzheimer cortex cDNA to mRNA,
                                                                                                                                                                                                                                                                                                                                              Homo
                                                                                 Unpublished 2 (bases 1
                                                                2 (bases 1 to 1832)
Isogai, T. and Yamamoto,
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                Fax:81-438-52-3986)
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                                  TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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Location/Qualifiers
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/codon_start=1
/protein_ic="BAC04368.1"
/protein_ic="BAC04368.1"
/db_xref="GI:21753576"
/translation="MWYLGIAATFCGLFLLPGFALQIOCYQCEEFQLNNDCSSPEFIV
/CTVNVQDMCQKEVMEQSAGIMYRSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPICNGPRPKKROSSASALRPGLRTTILFLKLALFSAHC"

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215. .640
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/tissue_type="alzheimer
/clone_lib="BRALZ2"
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/db_xref="taxon:9606"
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CCTTGCAGAGGACTGGCGCGGGACGCGAAGAGCAACGGGCGCTGCACAAAAGCGGCGCGCT 66
                                                      AATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACC
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                                         AATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACC
                                                                                             GACACGAGCAGGAGGAGAGACAGAGAAAAAGAAAACACAGCATGAGAACACAGTAAATA
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1 (bases 1 to 1890)
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/db_xref="GI:14272688"
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TPLCNGPRPKKRGSSASALRPGLRTTILFLKLASSRHTAELKEMPPPPALFFQPSPPT
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/db_xref="taxon:9606"
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                                                                                                                                                CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACCAATCAGCCC
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/db_xref="taxon:9606"
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Homo sapiens
                                                                                       Ota,T., Isogai,T., Nishikawa,T., Kawai,Y., Hayashi,K.
                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae. 1 (bases 1 to 587)
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                                                Secretory protein or membrane protein Patent: EP 1067182-A 620 10-JAN-2001; Helix Research Institute (JP)
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/db_xref="taxon:9606"
433 c 435 a ??
/organism="Homo sapiens"
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SYTLSCKLHTFLFEACSYATLLHVLTLSFERYIACHPERYKAVGPCQVKLLIGFVM
VTSALVALPLLFAMGTEYPLVNVPSHRGLTCNRSSTRHEQPETSNMSICTNLSSRWT
VFQSSLFGAFVYYLVVLLVLVSVAFMCWNMAQVLMKSQKGSLAGGTRPPQLAKKESEBESRT
ARRQTIIFLRLIVVTLAVCWMPNQIRRIMAAAKPKHDWTRSYFRAYMILLPFSETFFY
LSSVINPLLYTVSSQQFRRVFVQVLCCRLSLQHANHEKRLRVHAHSTTDSARFVQRPL
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/db_xref="GI:17901490"
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                    MCKee,K.K., Tan,C.P., Palyha,O.C., Liu,J., Feighner,S.D., McKee,K.K., Tan,C.P., Palyha,O.C., Liu,J., Feighner,S.D., Hreniuk,D.L., Smith,R.G., Howard,A.D. and Van der Ploeg,L.H. Cloning and characterization of two human G protein coupled receptor genes (GPR38 and GPR39) related to the growth hormone
                                                                                                                                                                                                                                                       Direct Submission Submitted (17-NOV
                                                                                                                                                                                                                                                                                                2 (bases 1 to 1362)
McKee, K.K., Tan, C.P.,
                                                                                                                                                                                                                                                                                                                                              secretagogue and neurotensin receptors
Genomics 46 (3), 426-434 (1997)
98110578
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                          hormone secretagogue
                                                                                                              /map="2q21-22"
/tissue_type="brain"
/dev_stage="fetal"
/product-"GPR39"
               /codon_start=1
                                        /note="orphan G
                                                         /gene="GPR39"
                                                                                   /gene="GPR39"
                                                                                                                                                           /chromosome="2"
                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
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Pred. No. 1.2e-207;
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Methods and compositions for the diagnosis and
Methods and compositions for the diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                 2528 bp
Sequence 3 from Patent WO0181634.
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Location/Qualifiers
                                                                                                                                                                                                                                   cardiovascular and tumorigenic dispatent: WO 0181634-A 3 01-NOV-2001
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Mammalia; Eutheria; Primates;
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                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
42. 1403
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LIGNSATIRVTGVLQKKGYLQKEVTDHAVSLACSDILVELIGHMEFYSIIWNEUTTS
SYTLSCKLHFFLFEACGSATLHAVILSFERY LAIGHBFRYKAVSGPGOVKLLIGEVW
VTSALVALPLLFAMGTEYPLVNVPSHRGLTCNRSSTRHHEQPETSNMSICTNLSSRWT
VFQSSLTGAFVVLVVLAVAFMCWNHMQVLMKSQKGSLAGGTREPPQLRKSESEBSRT
ARROTIFFLKLIVVTLAVAFWCWNHMQVLMKSQKGSLAGGTRAYMILLPESETEFY
ARROTIFFLKLIVVTLAVAFWCWNHMQVLMKSGKGLRGVAGAMHEKTFVARYMLYDSARFVQRPL
LSSVINPLLYTVSSQQFRRVFVQVLCCRLSLQHANHEKRLRVHAHSTTDSARFVQRPL
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LLGNSATIRVTQVLQKKGYLQKEVTDHMVSLACSDILVFLIGMPMEFYSIIWNPLTTS
SYTLSCKLHTFLFEACSYATLLHVLTLSFERYIAICHPFRYKAVSGPCQVKLLIGFVM
                                                    /protein_id="CAD19382.1"
/db_xref="GI:17901492"
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/db_xref="GI:2654161"
                                                                                        /codon_start=1
                                                                                                         /note="unnamed protein product"
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Pred. No. 1.2e-207;
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324 GCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAGGAGGGA 383
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                             AGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGG 323
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1 (bases 1 to 444)

Kroes, R. A., Moskal, J. R. and Yamamoto, H.

Differential gene expression in cancer
Patent: WO 0136685-A 95 25-MAY-2001;

NYXIS NeuroTherapies, Inc. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 95 from Patent WO0136685. AX150120
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/db_xref="taxon:9606"
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ARRQTIIFLRLIVVTLAVCWMENQIRRIMAAAKKHMDWRSYFBAYMILLPFSGFFFY
LSSVLMPLLYTVSSQQFRRVEYQVLCCRLSLQHANHEKRLRVBAHSTTDSARFVYDFI
LFASRRQSSARRTEKIFLSTFQSEAEPQSKSQSLSLESLEPNSGAKPANSAAENGFQE
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99.8%;
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| RESULT 12 ACC112072 LOCUS DEFINITION ACCESSION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      | Qy 564<br>Db 71                                    | Qy .504<br>Db 131                                                   | Oy 444<br>Db 191                                                 | Qy 384<br>Db 251                                                       | Db 311 |
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Submitted (17-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 13, 2002 this sequence version replaced g1:20303189.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (19-FEB-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA 3 (bases 1 to 92874)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Worley, K.C.
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished
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                                                                                                                                                                                                                                                                                                                                           NOTE: Estimated insert size may differ from sequence length
(see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a "working draft' sequence. It currently consists of 50 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.

This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                              as soon as it
be preserved.
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequencing vector: Plasmid; chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap; version 0.990329 consensus quality: 32512 bases at least Q40 Consensus quality: 34515 bases at least Q30 consensus quality: 35930 bases at least Q30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Center: Baylor College of Medicine Center code: BCM
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/clone="CH230-286021"
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MacCean, C., Macdonald, P., Major, J., Marguis, N., Matthews, C.,
MacCearthy, M., McEwan, P., McKernan, K., McPheeters, R., Meldrim, J.,
McCarthy, M., McEwan, P., McKernan, K., McPheeters, R., Meldrim, J.,
Meneus, L., Mihova, T., Mlenga, V., Murphy, T., Naylor, J., Nguyen, C.,
Norbu, C., Norman, C.H., O'Connor, T., O'Donnell, P., O'Neil, D.,
Oliver, J., Peterson, K., Phunkhang, P., Pierre, N., Pollara, V.,
Raymond, C., Retta, R., Rieback, M., Riley, R., Rse, C., Rogov, P.,
Roman, J., Rosetti, M., Roy, A., Santos, R., Schauer, S., Schupback, R.,
Seaman, S., Severy, P., Spencer, B., Stange-Thomann, N., Stojanovic, N.,
Strauss, N., Subramanian, A., Talamas, J., Tesfaye, S., Theodore, J.,
Topham, K., Travers, M., Travis, M., Trigilio, J., Vassiliev, H.,
Viel, R., Vo, A., Wilson, B., Wu, X., Wyman, D., Ye, W.J., Young, G.,
Zainoun, J., Zembek, L., Zimmer, A. and Zody, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Birren,B., Linton,L., Nusbaum,C., Lander,E., Ali,A., Allen,N., Anderson,S., Barna,N., Bastlen,V., BoguslavKiy,L., Boukhgalter,B., Brown,A., Camarata,J., Campopiano,A., Chang,J., Chazaro,B., Choepel,Y., Colangelo,M., Collins,S., Collymore,A., Cook,A., Cooke,P., DeArellano,K., Dewar,K., Diaz,J.S., Dodge,S., Faro,S., Certeratra,P., FittHugh,W., Gage,D., Galagan,J., Gardyaa,S., Cord,S., Cord
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AC099984 70395 bp DNA linear HTG 2 MUS musculus clone RP23-24D9, LOW-PASS SEQUENCE SAMPLING.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (22-NOV-2001) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA All repeats were identified using RepeatMasker: Smit, A.F.A. & Green, P. (1996-1997)
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Birren, B., Linton, L., Nusbaum, C. and Lander, E.
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                  sequencing reads that have not been assembled into contigs. Runs of N are used to separate the reads and the order in which they appear is completely arbitrary. Low-pass sequence sampling is useful for identifying clones that may be gene-rich and allows overlap relationships among clones to be deduced.
                                                                                                                                                                                                                                                                                                        NOTE: This record contains 84 individual
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                                                                                                                                                                                                                                                                                                                                                                                       Center clone name: 24_D_9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: sequence_submissions@genome.wi.mit.edu
------ Project Information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Web site: http://www-seq.wi.mit.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Center code: WIBR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Center: Whitehead Institute/ MIT Center for Genome Research
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19653 с
it should not be assumed that this clone
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will be sequenced to completion. In the event that the record is updated, the accession number will
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13431 13530: gap of 100 bp

13531 14266: contig of 736 bp in length

14267 14366: gap of 100 bp

14367 15105: contig of 739 bp in length

15106 15205: gap of 100 bp

15206 15323: contig of 727 bp in length

15933 16032: gap of
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0622: contig of 712 bp in length

6623 6722: gap of 100 bp

6723 7458: contig of 736 bp in length

7459 7558: gap of 100 bp

7559 8357: contig of 736 bp in length
27634 27733: gap of 100 bp
27734 28467: contig of 734 bp in length
28468 28567: gap of 100 bp
28568 29307: contig of 740 bp in length
29308 29407: gap of 100 bp
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8458 9202: contig of 745 bp in length
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1677 2451: contig of 775 bp in length
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18612: contig of 805

18612: gap of 100 h

19336: con+'-
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16767: contig of 735 bp in length

16867: gap of 100 bp

17607: contig of 740 bp in length

17707: gap of 100 bp

18512: contig of 805 bp in length
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26801: contig of 745 bp in
901: gap of 100 bn
27633: con++-
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21820: contig of 719 bp in length
120: gap of 100 bp
22647: contig of 727 bp in length
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19336: contig of 724 bp in length
136: gap of 100 bp
20162: contig of 726 bp in length
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25956: contig of 749 bp in length
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23424: contig of 677 bp in length
24: gap of 100 bp
24268: contig of 744 bp in length
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1250 30990: contig of 741 bp in length
1991 31090: gap of 100 bp
1991 31820: contig of 730 bp in length
121 31920: gap of 100 bp
121 32656: contig of 736 bp in length
157 32756: gap of 100 bp
167 32756: gap of 100 bp
178 33490: ---
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52922: contig of 728 bp in length
53022: gap of 100 bp
530755: contig of 733 bp in length
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17101: contig of 739 bp i
17101: gap of 100 bp
57848: contig of 747 hr
1948: gap of
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11307: contig of 699 bp 1
51307: gap of 100
52094: con**
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55385: contig of 715 bp
5485: gap of 100 bp
56162: contig of 677 bp
6262: gap of 100 bp
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54570: cont
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48723: contig of 737 bp in length
823: 100 bp
49570: contig of 747 bp in length
570: gap of 100 bp
50408: contig of 738 bp in length
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40227: contig of 718 bp in length
127: gap of 100 bp
41093: contig of 766 bp in length
193: gap of 100 bp
41952: contig of 759 bp in length
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ap of
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contig of 715 bp
                                    Score 24; DB 2;
Pred. No. 0.027;
         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished
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Submitted (24-MAR-2000) Multimegabase Sequencing Center, University Mashington, PO BOX 357730, Seattle, WA 98195, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98105, USA
On Nov 17, 2000 this sequence version replaced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rowen,L., Madan,A., Qin,S., Abbasi,N., Baradarani,L., Birditt,B., Bloom,S., Dors,M., Dickhoff,R., Fleetwood,P., Harrison,G., James,R., Kaur,A., Madan,A., Owen,M.P., Ratcliffe,A., Shaffer,T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (17-NOV-2000) Multimegabase Sequencing Center, Institute for Systems Biology, 4225 Roosevelt Way NE, Suite 200, Seattle, WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rowen,L., Madan,A., Qin,S., Baradarani,L., Birditt,B., Bloom,S., Dors,M., Dickhoff,R., Fleetwood,P., Harrison,G., Kaur,A., Madan,A., Nesbitt,R., Traicoff,R. and Hood,L.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: leerowen@systemsbiology.org
-------Summary Statistics
Sequencing vector: pUC18: L08752
Chemistry: Dye-terminator Big Dye; 90
Chemistry: Dye-primer Big Dye; 108 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Center: Multimegabase Sequencing Center Center code: UWMSC Web site: http://chroma.mbt.washington.edu/msg_www
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Assembly program: Phrap;
                                                                                                                                                                                                                                                                                                                                            /clone_lib~"Cal Tech human BAC library 11"
/note~"This clone overlaps RP11-313P18, RP11-414J1 and RP11-56B16. Data from all of these BACs were combined, with an attempt to generate the consensus sequence from CTD-2306G16."
                                                                                                                                                                    /note="Overlap with RP11-313P18, 66240. .72320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
1. .94718
'note="low quality data"
                                  /note="low quality data"
72960 .72980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="CTD-2308G16"
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                                                                                                                                     'note="number of Ts unclear"
                                                                                                                                                                                                                                                                note="overlap with RP11-414J1, AC020685"
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JOURNAL REFERENCE
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AUTHORS
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VERSION
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AC011639/c
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (08-OCT-1999) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA On May 4, 2001 this sequence version replaced gi:13123327. All repeats were identified using RepeatMasker: Smit, A.F.A. & Green, P. (1996-1997) http://ftp.genome.washington.edu/RM/RepeatMasker.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 167693)
Birren, B., Linton, L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                    Sequencing vector: M13; M7815; 12% of reads Sequencing vector: M13; M7815; 12% of reads Sequencing vector: Plasmid; n/a; 88% of reads Sequencing vector: Plasmid; bye; 100% of reads Assembly program: Phrap; version 0.960731

Consensus quality: 164534 bases at least Q40 Consensus quality: 165712 bases at least Q30 Consensus quality: 165712 bases at least Q30 Insert size: 170000; agarose fp Insert size: 16693; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (bases 1 to 167693)
                                                                                                                                                                                                                                                                                                                                                                    Contact: sequence_submissions@genome.wi.mit.edu
                                                                                                                                                                                                                                                                                                                                                                                                                            Center code: WIBR web site: http://www-seq.wi.mit.edu
                                                                                                                                                                                                                                                                                                            Center project name: L3327
Center clone name: 13_H_19
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88020. .88070
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coverage:
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Homo sapiens chromosome 15 clone RP11-762A1 map
                                                         AC022742.4 GI:13123383
HTG; HTGS_PHASE1; HTGS_I
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
                    Homo sapiens
                                     Homo sapiens.
                                                                                                                      SEQUENCE, 4 unordered pieces.
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NOTE: This is a 'working draft' sequence. It currently consists of 8 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will
                                                                                                                                                                                                                                                                                                                                                                                                                                        49583 a
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120448 167327: contig of 46780 bp in length
167228 167327: gap of 100 bp
167328 167693: contig of 366 bp in length.
Location/Qualifiers
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54882 54981: gap of 100 bp
54982 57393: contig of 2412 bp in length
57394 57493: gap of 100 bp
57494 64870: contig of 7377 bp in length
64871 64970: gap of 100 bp
64971 73018: contig of 8048 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      90140 90239: gap of 100 bp
90240 120347: contig of 30108 bp in length
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90240. .120347
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167328, .167693
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/clone="RP11-13H19"
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                                                                                                                                                                                                                                                                                                                                                           3.7%; Score 23;
100.0%; Pred. No.
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                                                            HTGS_DRAFT; HTGS_FULLTOP
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FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequencing vector: M13; M77815; 2% of reads
Sequencing vector: Plasmid; n/a; 98% of reads
Sequencing vector: Plasmid; n/a; 98% of reads
Chemistry: Dye-terminator Big Dye; 100% of reads
Assembly program: Phrap; version 0.960731
Consensus quality: 176032 bases at least Q40
Consensus quality: 176045 bases at least Q20
Consensus quality: 176615 bases at least Q20
Insert size: 186000; agarose-fp
Insert size: 176667; sum-of-contigs
Quality coverage: 9.0 in Q20 bases; agarose-fp
Quality coverage: 9.5 in Q20 base
* NOTE: This is a 'working draft' sequence. It currently
consists of 4 contigs. The true order of the pieces
is not known and their order in this sequence record is
arbitrary. Caps between the contigs are represented as
runs of N, but the exact sizes of the gaps are unknown.
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Submitted (06-FEB-2000) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA On Feb 25, 2001 this sequence version replaced gi:11990718.
All repeats were identified using RepeatMasker:
Smit, A.F.A. & Green, P. (1996-1997)
http://ftp.genome.washington.edu/RM/RepeatMasker.html
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------ Project Information
Center project name: L5677
Center clone name: 762_A_1
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                                                                                /chromosome="15"
/map="15"
/clone_lib="RPCI-11 Human Male BAC"
                                          /clone="RP11-762A1"
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                                                                                                                                                                                                                                                                                                                2621: gap of 100 bp
176967: contig of 64346 bp in length
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JOURNAL REFERENCE
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                                                                                                                                                                                                                                                                        Submitted (06-FEB-2000) Whitehead Institute/MIT Center for Genesarch, 320 Charles Street, Cambridge, MA 02141, USA 3 (bases 1 to 180837)
Birren, B., Nusbaum, C., Lander, E., Ali, A., Allen, N., Anderson, Barna, N., Bastien, V., Bloom, T., Boguslavkiy, L., Boukhgalter, I. Camarata, J., Chang, J., Chazaro, B., Choepel, Y., Collymore, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cooke, P., DeArellano, K., Dewar, K., Domino, M., Doyle, M., Fenestor, J., Ferreira, P., FitzHugh, W., Forrest, C., Gage, D., Galagan, J., Gardyna, S., Grant, G., Hagos, B., Heaford, A., Horton, L., Gardyna, S., Grant, G., Hagos, B., Heaford, A., Horton, L., Gardyna, S., Grant, G., Hagos, B., Heaford, A., Horton, L., Locke, K., Landers, T., Lehoczky, J., Levine, R., Lieu, C., Liu, G., Locke, K., Landers, T., Lehoczky, J., McBwan, P., McGurk, A., McKernan, K., McCheeters, R., Meldrim, J., Meneus, L., Morrow, J., Naylor, J., McPheeters, R., Meldrim, J., Meneus, L., Morrow, J., Naylor, J., Norman, C.H., O'Connor, T., O'Donnell, P., Olivar, T.M., Peterson, K., Pierre, N., Pisani, C., Pollara, V., Raymond, C., Riley, R., Rothman, N., Roy, A., Santos, R., Severy, P., Spencer, B., Stange-Thomann, N., Stojanovic, N., Subramanian, A., Talamas, J., Tesfaye, S., Theodore, J., Tirrell, A., Vassillev, H., Viel, R., Vo, A., Wu, X., Wyman, D., Ye, W.J., Simmer, A. and Zoff, M.
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Homo sapiens chromosome 15 clone RP11-81117 map 15,
SEQUENCE, 8 unordered pieces.
                                  Faro, S., Ferreira, P., FitzGerald, M., Gage, D., Galagan, J., Gardyna, S., Gord, S., Graham, L., Grand-Pierre, N., Hagos, B., Horton, L., Hulme, W., Iliev, I., Johnson, R., Jones, C., Kamat, Karatas, A., Kells, C., Landers, T., Levine, R., Lindblad-Toh, K. Liu, G., MacLean, C., Macdonald, P., Major, J., Matthews, C., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meldrim, J., Meneus, L., Mihova, T., Mlenga, V., McCarthy, M., Meneus, L., Mihova, T., Mlenga, V., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Matthews, C., Mat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Birren,B., Nusbaum,C. and Lander,E. Homo sapiens chromosome 15, clone RP11-81117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 180837)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zimmer, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished
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112622. .176967
/note="assembly_fragment"
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Naylor, J., Nguyen, C., Nicol, R., Norbu, C.,
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                         Lander,E., Ali,A., Allen,N., Anderson,S.,
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                                                                                                                       Lindblad-Toh, K.,
                                                                                                                                                                                                                                                                                                                                                                                                                 Anderson, S.,
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WORKING DRAFT
Norman, C.H.,
                                                                                                                                                         Kamat,A.,
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FEATURES
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JOURNAL
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Center clone name: 81.1.17

Center clone name: 81.1.17

Center clone name: 81.1.17

Sequencing vector: M13; M77815; 3% of reads
Sequencing vector: Plasmid; n/a; 97% of reads
Sequencing vector: Plasmid; n/a; 97% of reads
Sequencing vector: Plasmid; n/a; 97% of reads
Sequencing vector: Plasmid; n/a; 97% of reads
Sequencing vector: Plasmid; n/a; 100% of reads
Chemistry: Dy-terminator Big Dye; 100% of reads
Assembly program: Phrap; version 0.960731
Consensus quality: 177875 bases at least Q40
Consensus quality: 179781 bases at least Q20
Insert size: 180.200; agarose-fp
Insert size: 180.207; sum-of-contigs
Quality coverage: 7.8 in Q20 bases; agarose-fp
Quality coverage: 7.8 in Q20 bases; agarose-fp
Quality coverage: 7.8 in Q20 bases
* NOTE: This is a 'working draft' sequence. It currently
consists of 8 contigs. The true order of the pieces
* is not known and their order in this sequence record is
* arbitrary. Gaps between the contigs are represented as
* runs of N, but the exact sizes of the gaps are unknown.
* This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   http://ftp.genome.washington.edu/RM/RepeatMasker.html
------ Genome Center
Center: Whitehead Institute/ MIT Center for Genome Research
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             O'Connor,T., O'Donnell,P., O'Neil,D., Oliver,J., Peterson,K., Phunkhang,P., Pierre,N., Raymond,C., Retta,R., Rise,C., Rogov,P., Roman,J., Roy,A., Schauer,S., Schupback,R., Seaman,S., Severy,P., Smith,C., Spencer,B., Stange-Thomann,N., Stojanovic,N., Talamas,J., Tesfaye,S., Theodore,J., Topham,K., Travers,M., Vassiliev,H.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            208: contig of 208 bp in length
209 308: gap of 100 bp
309 6666: contig of 6358 bp in length
6667 6766: gap of 100 bp
6767 14021: contig of 7255 bp in length
14022 14121: gap of 100 bp
14122 21059: contig of 6938 bp in length
21160 21159: gap of 100 bp
21160 40187: contig of 19028 bp in length
40187: contig of 19028 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    be preserved.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: sequence_submissions@genome.wi.mit.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Center code: WIBR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100 bp

40288 68512: contig of 28225 bp in length

68513 68612: gap of 100 bp

68613 107892: contig of 500 bp

07893 107000
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107993 180837: contig of 72845 b
Location/Qualifiers
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R., Vo,A., Wilson,B., Wu,X., Wyman,D., Young,G., Zainoun,J.,
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                                                                                                                                vector_side:left"
                                                                                                                                                                    clone_end:SP6
/note-"assembly_fragment"
                                                                    /note=
                                                                                                       309.
                                                                                                                                                                                          /note="assembly_fragment
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                                                                                                                                                                                                                                                                                                                                                                             /chromosome="15"
                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
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RESULT 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Funke, R., Gage, D., Galagan, J., Gardyna, S., Gilbert, D., Grant, G., Hagos, B., Heaford, A., Horton, L., Howland, J.C., Jones, C., Kann, L., Karatas, A., Lehoczky, J., Lieu, C., Locke, K., Macdonald, P., Marquis, N., McEwan, P., McGurk, A., McKernan, K., McLaughlin, J., Meldrim, J., Molla, M., Morris, W., Morrow, J., Mychaleckyj, J., Naylor, J., Niloff, M., O'Connor, T., O'Donnell, P., Pavlin, B., Peterson, K., Pollara, V., Riley, R., Roberts, D., Roy, A., Severy, P., Peterson, K., Pollara, V., Riley, R., Roberts, D., Roy, A., Severy, P.,
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                                                                                                                                                                                                                              http://ftp.genome.washington.edu/RM/RepeatMasker.html
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                                               Center project name: L2273
Center clone name: 56_B_16
                                                                                                 ----- Project Information
                                                                                                                       Contact: sequence_submissions@genome.wi.mit.edu
Sequencing vector: M13; M77815;
                                                                                                                                                    Web site: http://www-seq.wi.mit.edu
                                                                                                                                                                            Center code: WIBR
                                                                                                                                                                                                      Center: Whitehead Institute/ MIT Center for Genome Research
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consists of 27 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence
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Insert size: 181885; sum-of-contigs
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36984 c 37799 g 54422
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23977. 27612
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18518. .20760
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141632. .163266
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31061. .34289
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27713. .30960
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107394. .124850
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On NOV 27, 2001 this sequence version replaced gi:13624393
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Direct Submission
Submitted (27-NOV-2001) Multimegabase Sequencing Center, Institute for Systems Biology, 4225 Roosevelt Way NE, Suite 200, Seattle, WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rowen,L., Madan,A., Qin,S., Baradarani,L., Birditt,B., Bloom,S. Burke,J., Dors,M., Fleetwood,P., Kaur,A., Madan,A., Nesbitt,R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (21-MAR-2001) Multimegabase Sequencing Center, Institute for Systems Biology, 4225 Roosevelt Way NE, Suite 200, Seattle, WA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rowen, L., Madan, A., Qin, S., Baradarani, L., Birditt, B., Bloom, S., Dors, M., Dickhoff, R., Fleetwood, P., Harrison, G., Kaur, A., Madan, A., Nesbitt, R., Traicoff, R. and Hood, L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequencing of human chromosome 15 D15S146-D15S117 region unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Center: Multimegabase Sequencing Center Center code: UWMSC Web site: http://chroma.mbt.washington.edu/msg_www
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: leerowen@systemsbiology.org
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                                                                      /note="overlap with CTD-2650P22 AC090970" 80765. .192464
                                                                                                                                                                                                                                                                    /note="Data from overlapping BACs CTD-2308G16, RP11-13H19, RP11-47K1, and CTD-2650P22 was added and the consensus sequence was determined from RP11-56B16 to the extent
                                                                                                                                                                                                                                               possible"
                                                                                                                                                                                                                                                                                                                                            /clone="RP11-56B16"
/clone_lib="RPCI human BAC library 11"
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/db_xref="taxon:9606"
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                                                                                                                                            note-"overlap with CTD-2308G16 AC026770"
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Best Local Similarity
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                                                                                                           Submitted (08-JUL-2002) Human Genome Sequencing Center, Depa of Molecular and Human Genetics, Baylor College of Medicine, Baylor Plaza, Houston, TX 77030, USA
                                         Direct Submission
                                                   Worley, K.C.
                                                                      Unpublished
                                                                                   Direct Submission
                                                                                             Wu,C., Wu,Y., Wu,Y.F., Zhou,J., Zorrilla,S., Nelson,D., Weinstock,G. and Gibbs,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AC126666
AC126666.1 GI:21703541
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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100.0%; Pred. No.
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*** SEQUENCING IN PROGRESS ***,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Worley,K.C.
Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NOTE: Estimated insert size may differ from sequence length
(see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).
NOTE: This is a 'working draft' sequence. It currently
consists of 61 contigs. The true order of the pieces
is not known and their order in this sequence record is
arbitrary. Gaps between the contigs are represented as
runs of N, but the exact sizes of the gaps are unknown.
This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap; version 0.990329
Consensus quality: 141149 bases at least Q40
Consensus quality: 153199 bases at least Q30
Consensus quality: 163387 bases at least Q20
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Center clone name: CH230-3P12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
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3823: contig
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100.0%; Pred. No.
                                                                                                                                        hap2a"
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Cheuk, R., Chen, H., Kim, C.J., Koesema, E., Meyers, M.C., Banh, J.,
Bowser, L., Carninci, P., Chang, E., Dale, J.M., Goldsmith, A.D.,
Bowser, L., Carninci, P., Chang, E., Dale, J.M., Goldsmith, A.D.,
Hayashizaki, Y., Ishida, J., Jones, T., Kamiya, A., Karlin-Neumann, G.,
Kawai, J., Lam, B., Lee, J.M., Lin, J., Miranda, M., Narusaka, M.,
Rquyen, M., Onodera, C.S., Palm, C.J., Quach, H. L., Sakurai, T.,
Satou, M., Seki, M., Southwick, A., Tang, C.C., Toriumi, M., Wu, H.C.,
Yamada, K., Yamamura, Y., Yu, G., Yu, S., Shinozaki, K., Davis, R.W.,
Theologis, A. and Ecker, J.R.
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Cheuk, R., Chen, H., Kim, C.J., Koesema, E., Meyers, M.C., Banh, J.,
Cheuk, R., Chen, H., Kim, C.J., Koesema, E., Meyers, M.C., Banh, J.,
Bowser, L., Carninci, P., Chang, E., Dale, J.M., Goldsmith, A.D.,
Hayashizaki, Y., Ishida, J., Jones, T., Kamiya, A., Karlin-Neumann, G.,
Kawai, J., Lam, B., Lee, J.M., Lin, J., Miranda, M., Narusaka, M.,
Nguyen, M., Onodera, C.S., Palm, C.J., Quach, H.L., Sakurai, T.,
Satou, M., Southwick, A., Tang, C.C., Toriumi, M., Wu, H.C.,
Yamada, K., Yamamura, Y., Yu, G., Yu, S., Shinozaki, K., Davis, R.W.,
work. Shinozaki,K. (RIKEN GSC) contributed equally to contributed equally to this work as PIs.

Location/Qualifiers

1. .1982
                                                                                                                                                                                                                  The Salk, Stanford, PGEC (SSP) Consortium members carried out the sequencing and annotation of the RAFL cDNAs: Cheuk,R., Chen,H., Kim,C.J., Koesema,E., Meyers,M.C., Shinn,P., Banh,J. Bowser,L., Dale,J.M., Goldsmith,A.D., Jiang,P.X., Jones,T., Karlin-Neumann,G. Lam,B., Lee,J.M., Lin,J., Liu,S.X., Miranda,M., Nguyen,M., onodera,C.S., Palm,C.J., Pham,P.K., Quach,H.L., Southwick,A., Tang,C.C., Toriumi,M., Yamada,K., Yamamura, Y., Yu,G., Yu,S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              RIKEN Genomic Sciences Center (GSC) members carried out the collection and clustering of RAFL CDNAS (RAFL CDNA: 'RIKEN Arabidopsis Full-Length cDNA'): Seki.M., Narusaka,M., Ishida,J., Satou,M., Kamiya,A., Sakurai,T., Carninci,P., Kawai,J., Hayashizaki,Y. and Shinozaki,K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Arabidopsis thaliana AT5g12840/T24H18_10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (30-Nov-2001) Salk Institute Genomic Analysis Laboratory (SIGNAL), Plant Biology Laboratory, The Salk Institute for Biological Studies, 10010 N. Torrey Pines Road, La Jolla, CA 92037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FLI_CDNA
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RTALPLDMAQEPVYVNAKQYEGILRRRKARAKAELERKVIRDRKPYLHESRHKHAMRR
ARASGGRFAKKSEVEAGEDAGGRDRERGSATNSSGSEQVETDSNETLNSSGAP"
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0.29;
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MEDLINE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Biologie Moleculaire des Plantes, Strasbourg 67084, FRANCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NtSET1, a member of a newly identified subgroup of plant SET-domain-containing proteins, is chromatin-associated and its ectopic overexpression inhibits tobacco plant growth plant J. 28 (4), 371-383 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (01-SEP-2000) Shen W.H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shen, W.H
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AJ294474.1
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                  /translation="MEGGLRSGGNNPPSIDKTRYLDVKPLRCLAPVFPSPNGMSSVST
PQPSPFVCVPPTGPFPSPVAPENSGRESSOOTPSGVPNOGGSPGFAOPI
SPVPLNSFRFTTTANGNSGRSRFAVDDDDYSNSODONDGFASGFSVHVNNVEDSGTGK
KRGRPKKPRRAQAEGLTPVEVDVEPLLTQLLTSFKLVDLDQVKRADGDKELAGRVLL
YFDLFRRAWTQIDESSBDFPGSGRRPDLKASNMLMTKGVRTNQTKRRIGNAPGIEVGDIF
FFRMELCLYGCLHAPTMAGIDYMSVKLITMDESPLAVSTRGVKDVAXPTGKTIYTVGDLYTYTGQ
GGVQRKDGQVFDQKLERGNLALEKSVHRANEVRVIRGVKDVAXPTGKTIYIVGLYKIQ
GSVQRKDGQVFDQKLERGNLALEKSVHRANEVRVIRGVKDVAXPTGKTIYIVGLXKIQ
GSVQRKDGQVFDQKLERGNLALEKSVHRANEVRVIRGVKDVAXPTGKTIXIVGLXGG
GGVQRCGQVFDQKLERGNLALEKSVHRANEVRVIRGVKDQXBRVGVILDDLTSGAES
QFVCLVNUVDDEKGFAYFTYIFSLKYSKFFVMKSJQQGFKARLEVFKTKRRGWGLR
SWDPIRGGGFICEYAGEVIDAGNYSDUNYIFDATRIYALEAERDYNDESRKVPFFLY
SVENCHUTSTANDAGNYSDUNYIFDATRIYALEAERDYNDESRKVPFFLY
                                                                                                                                                                                                                                                                                                                                                                                                                      /gene="set1"
157. .2271
                                                                                                                                                                                                                                                                                     /protein_id="CAC67503,
/db_xref="GI:15485584"
ISAKNGGNISRFMNHSCSPNVYWQLVVRQSNNEATYHIAFFAIRHIPPMQELTFDYGM
                                                                                                                                                                                                                                                                     /db_xref="SPTREMBL:Q93YF5"
                                                                                                                                                                                                                                                                                                            /product="SET-domain-containing protein"
/protein_id="CAC67503,1"
                                                                                                                                                                                                                                                                                                                                                         /codon_start=1
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/note="ecotype: Columbia"
/ 376 c 450 g 615 t
                                                                                                                                                                                                                                                                                                                                                                         function="chromatin modeling"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /cell_line="TBY2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="PBF83b"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /db_xref="taxon:4097"
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/db_xref="taxon:3702"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (24-NOV-1997) Entomology, Oklahoma State University, 127 Noble Research Center, Stillwater, OK 74078, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Palmer, M.J., Guo, > Direct Submission
                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ixodid tick, Amblyomma americanum (L.)
Mol. Cell. Endocrinol. 139 (1-2), 45-60 (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Guo,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amblyomma americanum retinoid AF035577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Isolation of two functional retinoid X receptor subtypes from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Acari;
Parasitiformes; Ixodida; Ixodidae; Amblyomma.
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                                                                                                                                                                                                                                                                              /Godon_start=1
/product="retinoid x receptor"
/protein_id="nac15588.1"
/protein_id="nac15588.1"
/db_xref="GI:3098334"
/translation="MATQERAPGINGGTSNGVSSSLPPOPSLPAERLRRALGSITVGR PAAQQRSSGIDTATAAATSHHALSTKEPLSGSKHLCSICGDRASGKHYGVYSCEGCKG FFKRTVRKDLSYACREERTCIIDKRQRNRCQVCRYQKCLACGMKRBAVQEERQRTKDR ADSEVESTGGGAPPEMPLERILEAELRVESQTGTLSESAQQODFYSSICQAADROLHQ
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AHGAGVGAIFDRVLTELVAKMREMKMDRTELGCLLAVVLFNPEAKGLRTCPSGGPEGE
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477 c 659 g 7
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DEFINITION

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The software programs used to predict genes include: Grail
(Informatics Group, Oak Ridge Mational Laboratory,
http://compbio.ornl.gov/Grail-1.3/),
GENSCAN (Chris Burge, MIT, http://CCR-081.mit.edu/GENSCAN.html),
MetGene2 (S.M. Hebsgdard, et al., CES, Technical University of
Denmark, http://www.cbs.dtu.dk/services/NetGene2/) and
SplicePredictor (Volker Brendel, Stanford University,
http://gremlinl.zool.iastate.edu/cgi-bin/sp.cgi).
http://gremlinl.zool.iastate.edu/cgi-bin/sp.cgi).
Genes encoding tRNAs are predicted by tRNAscan-SE
(Sean Eddy, Washington University School of Medicine, St. Louis,
http://genome.wustl.edu/eddy/tRNAscan-SE/
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Structural analysis of Arabidopsis thaliana chromosome 5.
Scruence features of the regions of 3,076,755 bp covered
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This sequence may not be the entire insert of this clone. It may be shorter because we remove overlaps between neighboring submissions. The 5' clone is MMN10 and the 3' clone is F15L12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 http://www.kazusa.or.jp/kaos/cgi-bin/agd_graph.cgi?c=MGO3
Genes with similarity to proteins in the databases are described in
'product' or 'note' qualifiers. Genes that have no significant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel:81-438-52-3935, Fax:81-438-52-3934)
Address for correspondence: kaos@kazusa.or.jp
For the latest information on annotation of this clone,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (29-OCT-1998) Yasukazu Nakamura, Kazusa DNA Research Institute, Department of Plant Gene Research; 1532-3, Yana, Kisarazu, Chiba 292-0812, Japan (E-mail:ynakamu@kazusa.or.]p,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence features of the regions pl and TAC clones pl and TAC clones plakes. 7 (1), 31-63 (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arabidopsis thaliana (strain:Columbia) DNA, clone_lib:Mitsui Pl clone:MGO3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AB019231.1 GI:3869070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AB019231 BA000015
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nakamura,Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20181125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (Sites)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (bases 1 to 43570)
                        1010(27)32. ...2799,3210. ...3339,3544. ...3693,3907. ...4081, 1011(27)32. ...2799,3210. ...3339,3544. ...3693,3907. ...4081, 270,4399. ...4592,4765. ...4874,4963. ...5062,5271. ...5483,5572. ...5720,6096. ...66237,6319. ...6459,6908. ...7096,7433. ...7680,7779. ...7872,8102. ...8336,8660. ...8851,9115. ...9213,9389. ...9592,9674. ...9866,10082. ...10273,10373. ...10524,10736. ...10891,10971. ...11081,11213. ...11307,11608. ...11812,11924. ...12126,12360. ...12411)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           complement(1224...210
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                                                                                                                                                                                                                                                              KNKDEDVVDEKEKEKLEDEQRSGERKKEKKKKRKSDEEIVSEERKSKKKRKSDEEMGS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /db_xref="taxon:3702"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1. .43570
                                                                                                                                                                                                                           EERKSKKKRKLKEIDD"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /codon_start=1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .2102)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    bp covered by sixty
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             please see
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CDS

VKGQSIDVVVELMKGHRSSLNMADSKGNTALHVATRKGRIKIVELLLDNNETSPSTKA

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EEHPLSGILGRUKGKGEFRAMUSKRVEFTGETVISDPNLKITEVOIPILMAQILI
FPECVSRHNIEKLRQCVRNGPNKYPGARNVRYDGSSRTLVGDYRKRIADELAIGCIV
FPECVSRHNIEKLRQCVRNGPNKYPGARNVRYDGSSRTLVGDYRKRIADELAIGCIV
DRHLQEGDVVLENRQPSLHRMSIMCHRARIMPRITLEFNESFLITEKDTETVDRAAFSL
LOSYMGDGMDSIDLPTPIILKPIELWTGKQIFSVLLRPNASIRVYVTLNVKEKNFKKG
EHGFDETMCINDGWYFRNSELISGOLGKATLGNGNKDGLYSTLLRDYNSHAAAVCMN
RLAKLSARWIGHIGFSIGIDDVOPGEELSKERKDSIOFGYDQCHRKIEEFNRGNLOLK
AGLDGAKSLEAETTGLHTIREATGKACMSGLHWRNSPLIMSQCGSKGSPINISOMX
CVGQQTVNGHRAPDGFIDRSLPHFPRMSKSPAAKGFVANSFYSGLTATEFFFHTMGGR
EGLVDTAVKTASTGYMSRRLMKALEDLLVHYDMTVRNASGCILOFTYGDDGMDPALME
EGLVDTAVKTASTGYMSRRLMKALEDLLVHYDMTVRNASGCILOFTYGDDGMDPALME
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PLTYVLDPNLVLGLFKRMSDKDCELLYIAYRPENLIITCMLVPPLSIRPSVMIGGIQS
NENDLTARLKQIILGNASLHKILSQPTSSPKNMQVWDTVQIEVARYINSEVRGCQNQP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GKDGAPLNENRLELKVQATCPPRSHHTYLSSEELSQKEBEELVRHDKSRVCTDAFVKS
LREFVSLLGVKSASPPQVLYKASGQKYEILPLIYINMQVFVKICVFRYREKKIBAGTA
IGTIGAQSIGBPGTQMTLKTFHFAGVASMNITQGVPRINBIINASKNISTPVISABLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /translation="metkmeleftkkpyleDvGpLkiksInfsvLsDLevmkAAEVQVwnIGLYDHSfkpyeNgLLDDpvNNGppNkKsICTTCEGNFQNCPGHYGYLKLDLPVNNVG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               complement(join(12822. .13766,13852. .14406))
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EAARTTIIDEIGTVMGNHGMSIDIRHMMLLADVMTYRGEVLGIQRTGIQKMDKSVLMO
ASFEBTGDHLFSAAASGKVDNIEGVTECVIMGIPMKLGTGILKVLQRTDDLPKLKYGP
DPIIS"
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VKNSILKTPRIKLNDNDIRVLDTGLDITPVVDKSRAHFNLHNLKNVLPNIIVNGIKTV
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product "ankyrin-like protein"
product "ankyrin-like protein"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /translation="mdsssllpsqwsdlpldilelisdrldhdssdtihllclrsvca
Twrlslplsnknnrlskfpkylpfwsssssssgfftlkqsnvykleaplnprtclvkl
QETTPGIMRVLDLfsndrictlpenpfskidlqefhyklvrrtykmeyanngggavpc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /evidence=not_experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene_id:MGO3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                complement(17742. .18866)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNMLWMYDILIEKKLCSEFYKLWADQKELANLHSKIPTNYRHEISKITAQICVGIGKG
RILVNRETREAVLNTWLEALYDDFGWMRRLSSRSLDRKLVEDGLSQTILTLSLRQQOV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLEELHLPDDSVTLILQRVSSQPSTSSTRTRTDDIFSKLLTGVLQAKDDKARREMKVL
                                                                                                                                                                                                                                                                                                                                                                                                                     YSDYRNGGRSRGICSDGDGVFNVDMQGDFVFSIKPNYFGP" complement(join(19582..20451,20673..210)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ILMKWFDRFLSKGDDCPNVQRAFEVWWRRAFIRQVLTEPDEPRLQITLY"
                                                                                                                                                                                                                                                                                                                                                                              /note="gene_id:MGO3.5"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CKSDISKSVYEFRVYNLKREEKRWEEVRDLGDVALFIGDDCSFSVQNPAGDLAGGFIF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNRCIVVDDKGKTVIYDVDFKVSDLAEGLVGGGGHKKHLVECSGGEVFLVDKYVKTVW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FWSLHSDKVVILSSGEDSAIIAIHSGGKLGFLKSGNDEKWKILDNSWNVIYEDIMLYK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /codon_start=1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IFKLVREEADYDVSRDTLYGLCHRCLTSLVLCLSEVTTQMNDPGKDRGALMGEIAREA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /codon_start=1
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CDS

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ACCESSION
VERSION
KEYWORDS
                                                                                                                                                                                                                                             RESULT 27
AC114102
                                                        SOURCE
                                                                                                                                                                                          DEFINITION
                                                                                                                                                                                                                         LOCUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                               ORGANISM
                                                                                                                                                                                                                                                                                                                                                        33922 GAATAAAACCATAAAATATTTA 33901
                                                                                                                                                                                                                                                                                                                                                                                                              426 GAATAAAACCATAAAATATTTA 447
                                                                               AC114102
AC114102.2 GI:21732042
HTG; HTGS_PHASE1.
                                                                                                                                                                                          Rattus norvegicus clone CH230-70H23,
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                               Rattus norvegicus
                                                  Norway rat.
                                                                                                                                                                                                                         AC114102
                                                                                                                                                                   ***, 56 unordered
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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IMHRIEASNUFKSKKESMARSROSGLLEFSGILTKRYKAI
IMHRIEASNUFKSRKESMARSROSGLEEFSGILTKRYKAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GAVDFLVKPIRKNELKNLWQHVWRRCHSSSGSGSESGIHDKKSVKPESTQGSENDASI
SDEHRNESGSSGGLSNQDGGSDNGSGTQSSWTKRASDTKSTSPSNQFPDAPNKKGTYE
NGCAHVNRLKEAEDQKEQIGTGSQTGWSMSKKAEEPGDLEKNAKYSVQALERNNDDTL
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LEDEKSCIDLVLTEVDMPVHSGTGLLSKIMSHKTLKNIPVIMMSSHDSMVLVFKCLSN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IYEFMANGDLERHIFAHKTMVEGPRPKGLTLPIRVLDWYTRMKIAVGVAKGLVYLHEE
LKVINADVKAGNILLDADFVPKLTDFGLATKIYEDENGYEQOGNIIPMKASMGYIAPE
GEISRLYSTRIDVYSYGAFLLVLFTGRKPFYSDNPAGKKULTDWETLRVADRIEDADVK
ADVALGNRYSVEGLKKIFDTARMCIKAERLERPTMSEVEAMVLEAAAFPVQVPRQVTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        complement(join(25224. .25460,25563. .25996,26073. .26262,
26345. .26514,26702. .26867))
/notce-"contains similarity to protein kinase
gene_id:MGO3.7"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VLLTGRKAYDLARPVAKEKLTDWLMPVWTRLEYAPMIVDVALGNKYSIEGLNRLLQTARMCINPQALERPAMDEVETMVREAAAFPVPKEPLVKERISTSK"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /translation="msnrflcclggggsskymgdpselpggggpggsdssdspdpgggs

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AHARQYQGLTLDTRVLDMNTRLRIAVGYAEGLVYVHQGLYAIHRDIKVERBILLDKDFV

PKLTDEGFATKIVYNSDGVERLREFNTRGTQGYIAPEADEFALVSTKSDIYSYGVLLL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HLRLPDDREVADPTAALLPVSPPLPDNCGESFLWRELVNGTINFRDEFYLGKGNFGEV
HRCYFSRLNVDGAVKVQNPQNPTGHVEFLAEVTTLRATNHPNVIRLLGNCYGQRNRAI
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/protein_id="BAA96937.1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     complement(join(22323.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                similar to unknown protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RRSAST"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /evidence≖not_experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /codon_start-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /codon_start=1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .23541,23892.
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Direct Submission
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Submitted (07-MAR-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
3 (bases 1 to 86093)
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Buhay, C., Burcell, K.L., Byrd, N.C., Carron, T.F., Carter, M., Cavazos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Davila, M.L., Davis, C., Day-Carroll, L., Dederich, D.A., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwalte, K. J., Draper, H., Dugan-Rocha, S., Durbh, K.J., Davis, D., Dugan-Rocha, S., Durbh, K.J., Davis, D., Dugan-Rocha, S., Durbh, K.J., Davis, D., Davis, D., Dugan-Rocha, S., Durbh, K.J., Davis, D., Davis, D., Dugan-Rocha, S., Durbh, K.J., Davis, D., Davis, D., Dugan-Rocha, S., Durbh, K.J., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis, D., Davis
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Submitted (13-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
On Jul 11, 2002 this sequence version replaced gi:19224429.
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Sequencing vector: Plasmid; Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap; version 0.990329 Consensus quality: 10778 bases at least 040 Consensus quality: 11488 bases at least 030 Consensus quality: 12089 bases at least 020
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NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 56 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
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2 (bases 1
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AL353013
AL353013.1
                                                   Arabidopsis thaliana.
Arabidopsis thaliana
Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots;
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots;
Spermatophyta; Magnollophyta; euroside; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                     Bevan, M., Robben, J., Mewes, H.W., Rudd, S.,
                                                                                                                                                        Arabidopsis
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                                           (bases 1 to 90020)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (19-APR-2000) MIPS, at the Max-Planck-Institut fuer Biochemie, Am Klopferspitz 18a, D-82152 Martinsried, FRG, E-mail: lemcke@mips.biochem.mpg.dee/mayer@mips.biochem.mpg.de Project Coordinator: Mike Bevan, Molecular Genetics Department, Cambridge Laboratory, John Innes Centre, Colney Lane, NR4 7UJ Norwich, UK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Laboratory, John Innes Centre, Col E-mail: michael.bevan@bbsrc.ac.uk
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complement()oin(2964...3167,3349...)
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4048...4148.4502...4778))
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/codon_start=1
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8885. .11nns
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /translation="mqskpgreneeevnnhhavqqpmmyaepwknnsfgvvpqarps GIPSNSSSLDCPNGSESNDVHSASEDGALNGENDGTWKDSQAATSSRSVDNHGMEGND FALSIRNMHDQFLVQPPELVGHYIACVENPYQDDYYGGLMGAYGHQQLGFRPYLGMPR ERTALPLDMAQEPYVNAKQYEGILRRRKARAKAELERKVIRDRKPYLHESRHKHAMR RARASGGRFAKKSEVBAGEDAGGRDRERGSATNSSGSEQVETDSNETLNSSGAP" complement(2964...3167)
                                                                                                                                                                                     complement(4502. .4778)
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/variety-"Columbia"
/db_xref="taxon:3702"
/chromosome="5"
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/note="similarity to zinc finger transcription
(PEII), Arabidopsis thaliana, EMBL:AF050463"
                                                                                                                                                                                                                                                                               complement(4149
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/TIANSIA tion="MASIALSGSCSLAFPLKSRSLSLPRPPSSSLNLTKPLRSLDSRF
SILKSPLEVSLRRRSSTLVKASSTVASASSSTPPLVPAPVPMGGAAIKPLLASIATG
LILMFVPVPEGVTRNAWQLLAIFLATIVGIITQPLPLGAVALMGLGASVLTKTLTFAA
AFSAFGDPIPMLIALAFFFARGFIKTGLGNRVAYOFVRLEGSSSLGLEVSLVFSEALL
APAIPSVSARAGGIFLPLVKSLCVACGSNVGDGTEHRLGSWLMLTCFQTSVISSSMFL
TRAMANDLSANLAFNTIKOTIGWTDWAKAAIVPGLUXLIVVPFLLXVLIYPPTVKSSPD
APKLAQEKLDKMGPMSKNELIMAATIFLTVGLIWIFGAKLGVDAVTAAILGLSVLVTG
VTWKECLAESVAWDTLTWFAALIAMAGYLNKYGLIEWFSQTVVKTVGGLGLSWQLSF
GILVLLYFYTHYFFAASGAAHIGAMFTAFLSVSTALGTPYFFAALVLAFLSNLMGGLTH
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PGSPSAAQHSTTPFISPSGNGSMPHSSMGWPQQNIPALNLPGSNIQLGRLRSSLNARD
IPSEQLSMLHEFEMQRDLAGDMHSPREMNHSARRWTLNDSNLEELFSACASPRFSDQ
LAVSSYLSPSHRSALLNQLQNIKQSMLSPIKTNLMSSPKNVEQHSLLQACASPRGGEP
ISPMNARMKQQLHSRSLSSRDFGSSSLPRDLMPTDSGSPLSPMSSWDQTHGSKVDWSVQ
SDELGRLRKSHSLANNPNREADVSWAQQMLKDSSSPRNGNRVVNMNGARPLTQGGSSV
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VSSPTKPHGTDVTFASEKKEYPIDPSLPDIKSGIYSTDEFRMFSFKIRPCSRAYSHDW
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DVAKNAGLQRCGKSCRLRWINYLRPDLKRGAFSPQEEDLIIRFHSILGNRWSQIAARL
PGRTDNEIKNFWNSTIKKRLKKMSDTSNLINNSSSSPNTASDSSSNSASSLDIKDIIG
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/gene="T24H18_30"
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ILSFPEAELNLSCGPDKSTALHCAASGASVNSLDVVKLLLSVGADPNIPDAHGNRPVD
                                                                                                                                                                                                                                                 /note="Contains ATP/GTP-binding site motif A (P-loop)
AA51-58; Myb DNA-binding domain repeat signatures
AA23-31; Myb DNA-binding domain repeat signatures AA95
                                                                                                                                                                                                                                                                                                                                          complement(join(14756. .15317,16476. .16756))
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                                                                                                                                                                                                                                                                                                                                                                                                        complement(join(14756. .15317,16476. .16756))
/gene="T24H18_40"
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/gene="T24H18_30"
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/gene="T24H18_30"
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SFMSLQEQGFVNPSLTHIQTNNPFPTGNMISHPCNDDFTPYVDGIYGVNAGVQGELYF
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                                                                                                                   /product="putative transcription factor (MYB46)"
/protein_id="CAB88251.1"
/db_xref="GI:7630043"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /gene="T24H18_40"
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Best Local Similarity
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Muzny, D.M., Adams, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbaria, J., Benton, J., Binage, K., Blankenburg, K., Bonnin, D., Bouck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Burkett, C., Burrell, K.L., Byrd, N.C., Carron, T.F., Carrter, M., Cavazoo, S.R., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Davila, M.L., Davis, C., Davy-Carroll, L., Dederich, D.A., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J.,
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***, 50
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complement(21475. .22941)
/gene="T24H18_60"
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complement(20531.
/gene="T24H18_50"
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/gene="T24H18_50"
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complement(20531. . 20752)
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/db_xref="GI:7630044"
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Homsi,F., Howard,S., Huber,J., Hulyk,S., Hume,J., Jackson,L.E.,
Jacobson,B., Jla,Y., Johnson,R., Jolivet,S., Joudah,S.,
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Nguyen,N., Nickerson,J., Newtson,N., Nguyen,A., Nguyen,N.,
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Nguyen,N., Nguyen,N., Nguyen,
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                                                                                                                                                                                                                                * (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

* NOTE: This is a 'working draft' sequence. It currently

* consists of 50 contigs. The true order of the places

* is not known and their order in this sequence record is

* arbitrary. Gaps between the contigs are represented as

* runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (10-APR-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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Falls,T., Ferraguto,D., Flagg,N., Ford,J., Foster,P., Frantz,P.,
Gabisi,A., Gao,J., Garcia,A., Garner,T., Garza,N., Gill,R.,
Gorrell,J.H., Guevara,W., Gunaratne,P., Hale,S., Hamilton,K.,
Harris,C., Harris,K., Hart,M., Havlak,P., Hawes,A., Hernandez,J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               * NOTE: Estimated insert size may differ from sequence length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Worley, K.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Direct Submission
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                                                                                                                                     as soon as it is available and the accession number will be preserved.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap; version 0.990339 Consensus quality: 4565 bases at least Q40 Consensus quality: 49416 bases at least Q30 Consensus quality: 51546 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequencing vector: Plasmid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Center project name: GTRW Center clone name: CH230-49C23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Center code: BCM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Center: Baylor College of Medicine
         \frac{1036}{1136}
1035: contig of 1035 bp in length 1135: gap of unknown length 2169: contig of 1034 bp in length
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COMMENT

TITLE

TITLE

2170 2270 2270 3323 3423 4694 4794 5881 5981

gap of contig gap of contig

unknown of 1271 of 1053

bp in 1

length

contig gap of contig gap of contig

unknown unknown

length

length

bp in length

length

length length

7413 7513 8600 8700 10029 10129 111471 111571 111571 13102 14816 14816

gap of contig gap of contig gap of contig gap of contig gap of contig gap of

of 1361 unknown of 1614 unknown

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unknown of 1638 unknown of 1115

contig

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unknown of 1056

length

bp in

10128 10028:

gap of contig gap of contig

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in length

unknown of 1531 unknown of 1342

ength

length length

length

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RESULT 30
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                                                                                                                                               Direct Submission
Submitted (28 JUN-2001) IBI/MIPS, National Research Center for Environment and Health, Ingolstaedter Landstr. 1, 85758 Neuherberg, FRG, E-mail: schoof@gsf.de,kmayer@gsf.de Project Coordinator: Mike Bevan, Molecular Genetics Department, Cambridge Laboratory, John Innes Centre, Colney Lane, NR4 7UJ Norwich, UK, E-mail: michael.bevan@bbsrc.ac.uk
                                                                             Information on performance of analysis and a more detailed annotation of this entry and other sequences of chromosomes 3, 4 and 5 can be viewed at: http://www.mips.biochem.mpg.de/proj/thal/
                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 98506)
Bevan, M., Murphy, G., Ridley, P., Hudson, S.,
Rudd, S., Schoof, H. and Mayer, K.F.X.
                                                                                                                                                                                                                                                                                                                  EU Arabidopsis sequencing, project.
                                                                                                                                                                                                                                                                                                                                                                       Rudd,S., Sci
Unpublished
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Arabidopsis thaliana
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/variety="Columbia"
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                                                                    Location/Qualifiers
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/clone="CH230-49C23"
20209 c 20180 g 22
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94029: contig of 1780
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92149: contig of 3153
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|----------------------------------------------------------|----------------------------------------------------------|-----------------------------------------------------|----------------------------------------------------------|----------------------------------------------------------|-------------------------------------|--------------------------------------------------------|----------------------------------|------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------|------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------|------------------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------|-------------------------------------------------------------------|------------------------------------------------------------------------------------------------|---------------------|-------------|--------|----------------------------------------------------------|
| /number=3<br>complement(1537315451)<br>/gene="At5g12280" | complement (1526515372) /gene="At5g12280"                | complement(19142. 19204) /gene_At5g12280" /nimhor=3 |                                                          | /gene="At0914280"<br>/number=2<br>complement(1507015141) | /number=1<br>complement(1439815069) | complement(1433214397)<br>/gene="At5g12280"            | /gene="At5g12280" /number=1      | QFLAQHPGSSTIMVSVPDPDDWKVVKITVQSLSENVASLKEKISGEIQFPTNKQKLRG<br>KAGFLKDNTSLAHYNVGAGEILFLSWW"<br>COMDINGENTTIAOGS 114371) | DYVLPBGYTIEDYDTIELTAQIYGRYGBEEWLDLIKEYDHKROGEELRPADSKEDYEN<br>RLSVVASEGIKRSEKLTSSRMAIVSKAFEYHLRRONEKRELTHKEGAAKACTLFGALY<br>AFYGPDGCFADIKDEDLPLLQHPSAIKWILTKLOPRMPPLGGESEEPEFDESVLVPED | /translation="MVTTLSpDTPDTQFISLCYKLQIVEVVRSPLFSTSRPGGSRGRA<br>/translation="MVTTLSpDTPDTQFISLCYKLQIVEVVRSPLFSTSRPGGSRGRA<br>MRSRLRLHKPIILGQMKFGESDFDVSYDLIFMLEKDNSNEAPAEPPREMRIHMDKIAR<br>VVARKGLRIERKIMNVSEYLAKNQDEGAQEPEAPATLKGDAQHYRRPFLLSDPAPKFP |                                                          | <pre>/note="similarity to splicing factor SF3a 120K chain,<br/>human, PIR:S60735"<br/>/codon_start=1</pre> | complement(join(1409514331,1439815069,1514215264,<br>1537315461,1556915717))<br>/gene=Mat5q12280" | .14331,1439815069,15142<br>15717))                                                                                                                                                              |                                                                                                                           | complement(1288613382)<br>/gene="At5g12270"<br>/number=2                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | /gene="At5g12270"<br>/gene="At5g12270"<br>/number=1                                                                                                                                    |                                                                                   | VYSDYLKFFEGRPHDGKTIESIKLP"  complement(12212. 12797) | WPSVCKEQVIDHFKRIKPITERILAILINNLNTIIDESNKETLMGTMRMNENYYPKCP<br>EPSLAIGTGHHSDINTLTLLLQEDGVLSSLYARAFEDGDKWIHVPPIPGAIVVNIGDV<br>IOTI SUNDAVESVHENVYVNKRYCSRVSIDVYCGDVHDSVIEDIDEVLDKNNEMARYRKI | FTSKILLGSPYVIDVSKNUEPHVAREICHARSKIGLFQIVNHGIAPAEFKGVIAAA FTSKILLGSPYVIDVSKNUEPHVAREICHARSKIGLFQIVNHGIAPAEFKGVIAAA RGFFELPAEERRYWRGSSVSETAWLTTSFNPCIESVLEWRDFLKFEYLPQRHDFAAT | /protein_id="cacazoo.i"<br>/db_xref="gI:14586357"<br>/*->                                      | /product-"1-AMINOCYCLOPROPANE-1-CARBOXYLATE OXIDASE-like protein" | /notem="Simitatity to ripening protein bo, commet, PIR:S01642" /codon start=1                  | 221212/9/,120001330 | 12797,12886 |        | e f                                                      |
|                                                          |                                                          | exon                                                | intron                                                   | exon                                                     | דוורדטוו                            |                                                        | exon                             | intron                                                                                                                 | exon                                                                                                                                                                                   | intron                                                                                                                                                                                                                                               | exon                                                     | intron                                                                                                     | exon                                                                                              |                                                                                                                                                                                                 |                                                                                                                           |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                                                                                                                                                                        |                                                                                   |                                                      |                                                                                                                                                                                           |                                                                                                                                                                             | CDS                                                                                            |                                                                   | gene                                                                                           | gene                |             |        | intron                                                   |
| /gene="At5g12290"<br>/number=6                           | /gene="At5g12290"<br>/number=6<br>complement(1758617808) | /number=5<br>complement(1751317585)                 | /number=0<br>complement(1743417512)<br>/gene="At5q12290" | complement(1740517433)<br>/gene="At5912290"              | /gene="At5g12290"<br>/number=4      | /gene="At512290"<br>/number=4<br>/number=117319 17404) | /number=3 complement(1719517318) | /number=3<br>complement(1705417194)<br>/gene="At5g12290"                                                               | /number=/<br>complement(1694317053)<br>/gene="At5912290"                                                                                                                               | /number=2<br>complement(1682616942)<br>/gene="At5g12290"                                                                                                                                                                                             | /number=1<br>complement(1672516825)<br>/gene="At5g12290" | /number=1<br>complement(1657416724)<br>/gene="At5g12290"                                                   | NSUTITIONS                                                                                        | VERNEU VIRAKLALDAFFES IVMLTVLRTWLKKDSKAQGRGRIARIHRRLLVVEIEKRIMQYQS<br>NFAILAALDAFFUS IVMLTVLRTWLKKDSKAQGRGRIARIHRRLLVVEIEKRIMQYQS<br>YIEQGRDKDAETVFGLLIYSLERLYRVVEKPARATDEWDFETRSDRVRKATAANIIQA | PEOPILS IRDELFOT FRKRHKGVMETEEVOLTOOSLIRMLRNECEQATREKVPDNASD OPNITYVMNDVDKET, URDTHNIJSGELARGILIOVOKIKIDIETAMLELDOILRANEI | SNIHHIIQAS DIE MANAGEGSTAMAAR IN IN SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SOOD ON SO | /translation="MDSQPPVNESTPSTALSNESELVPFVSSYLWNKLASLLFTSKr1 FLGKISNLYRQTVSRKRSISFPLFLDFSDFPSSSTITSNVSADTARIHGVLEEIMADVL FLGKISNLYRQTVSRKRSISFPLBADGAAVENTEEBGDFAAFVNESTKFVSKSISFDSANGHL | /product="putative protein"<br>/protein_id="AcA42880.1"<br>/db_xref="GI:14586359" | AA305-328"<br>/codon_start=1                         | /gene ncygizzy /note= ncygizzy /note= nsimilarity to NCA2 protein, yeast, PIR:S54389 /noteins 'Homeobox' domain signature and profile                                                     | 3659,1883018946,19038.<br>3719,1991520147))                                                                                                                                 | complement(join(1648616573,1672516825,1694317053, 1719517318,1740517433,1751317855,1780917898, | 3719,1991520147))                                                 | complement(join(1648616573,1672516825,1694317053, 1719517318,1740517433,17513175717809. 17188, |                     | 280"        | / your | /number=4<br>complement(1545215568)<br>/dene="xt5d12280" |

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RESULT 31
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Li,J., Li,Z., Lichtarge, U., Lieu, U., Lina,R., Ma,J., Lozado,R.J., Lu,X., Lucter,A., Lucter,R., Luna,R., Martinez,E., Mahashwari,M., Mapua,P., Martin,R., Martindale,A., Martinez,E., Maksbwari,M., Mei,G., Metzker,M., Massey,E., Mawhiney,E., McLeod,M.P., Meador,M., Mei,G., Metzker,M., Massey,E., Mawhiney,E., McLeod,M.P., Meador,M., Mei,G., McTris,S.,
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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/gene="At5g12290"
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/gene="At5g12290"
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/gene="At5g12290"
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/gene="At5g12290"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (24-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (07-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    * NOTE: Estimated insert size may differ from sequence length
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NOTE: This is a 'working draft' sequence. It currently consists of 50 contigs. The true order of the pieces is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            as soon as it is available and the accession number will be preserved.
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Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 5595 bases at least Q40
Consensus quality: 60532 bases at least Q40
Consensus quality: 633142 bases at least Q20
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Center clone name: CH230-303E15
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Project Information
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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, *** SEQUENCING IN PROGRESS ***,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (12-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, Ty 77030, USA
On Jul 11, 2002 this sequence version replaced gi:18138579.

Center: Baylor College of Medicine
Center: Baylor College of Medicine
Center code: BCM
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Direct Submission
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NOTE: This is a 'working draft' sequence. It currently consists of 171 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as
                                                                                                                                                                                                                                                                                                                                                           arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will
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Submitted (12-JUL-2002) Human Genome Sequencing Center, Depar
of Molecular and Human Genetics, Baylor College of Medicine,
Baylor Plaza, Houston, TX 77030, USA
On Jul 11, 2002 this sequence version replaced gi:17973362.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       regions were either double-stranded or sequenced with an alternate chemistry or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by at least one plasmid subclone or more than one M13 subclone; and the assembly was confirmed by restriction digest. The following abbreviations are used to associate primary accession numbers given in the feature table with their source databases: Em:, EMBL; Sw:, SWISSPROT; TT:, TREMBL; WP:, WORMPEP; Information on the WORMPEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     humquery@sanger.ac.uk Clone requests: clonerequest@sanger.ac.uk on Apr 10, 2002 this sequence version replaced gi:19068255.

During sequence assembly data is compared from overlapping clones. Where differences are found these are annotated as variations together with a note of the overlapping clone name. Note that the variation annotation may not be found in the sequence submission corresponding to the overlapping clone, as we submit sequences with only a small overlap as described above.

This sequence was finished as follows unless otherwise noted: all the corresponding to the overlapping clone, as we submit sequences with only a small overlap as described above.
                                                                                                                                                                                                                                                                                                                                  http://www.sanger.ac.uk/Projects/C_elegans/wormpep RP23-243J8 is from the RPCI-23 Mouse PAC Library constructed by the group of Pleter de Jong. For further details see http://www.chori.org/bacpac/home.htm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (21-MAY-2002) Wellcome Trust Sanger Institute, Hinxton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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/chromosome="11"
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                                                                                                 /clone_lib="RPCI-23"
27406 c 26381 q 2
                                                                                                                                                      /clone="RP23-243J8"
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contig of 1198 bp in length
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contig of 1855 bp in length
Score 22; pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1SA, UK. E-mail enquiries:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 22;
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NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 75 contigs. The true order of the pieces is not known and their order in this sequence record is arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown.

This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequencing vector: Plismid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 34683 bases at least 040
Consensus quality: 37389 bases at least Q30
Consensus quality: 37389 bases at least Q20
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REFERENCE AUTHORS

(bases 1 to 115643)

SOURCE

ORGANISM

Rattus norvegicus

VERSION

AC094168 AC094168.3 GI:21700340

HTGS\_PHASE1.

KEYWORDS ACCESSION DEFINITION

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RES MUZNY, D.M., Adams,C., Adlo-Oduola,B., Ali-osman,F.R., Allen,C., Alsbrooks,S.L., Amaratunge,H.C., Are,J.R., Ayele,M., Banks,T., Alsbrooks,S.L., Amaratunge,H.C., Are,J.R., Ayele,M., Banks,T., Barbooks,S.L., Benton,J., Binage,K., Blanksabburg,K., Bontin,D., Bouck,J., Bovie,S., Brieva,M., Brown,E., Brown,M., Bryant,N.P., Bouck,J., Chen,G., Chen,G., Chen,G., Chen,G., Chen,G., Chen,R., Chen,Z., Chowdhry,I., Christopoulos,C., Cleveland,C.D., Cox,C., Coyle,M.D., Dathorne,S.R., David,R., Devila,M.L., Davis,C., Davy-Carroll,L., Dederich,D.A., Davila,R., Delaney,K.R., Delgado,O., Denn,A.L., Ding,Y., Durbin,K.J., Douthwaite,K.J., Draper,H., Dugan-Rocha,S., Durbin,K.J., Delaney,K.R., Delgado,O., Denn,A.L., Ding,Y., Durbin,K.J., Carcia,A., Garner,T., Gerza,N., Gill,R., Gabisi,A., Gao,J., Garcia,A., Garner,T., Gerza,N., Gill,R., Gabisi,A., Gao,J., Garcia,A., Garner,T., Gerza,N., Gill,R., Gorrell,J.H., Guevara,W., Gunaratne,P., Hale,S., Hamilton,K., Gorrell,J.H., Guevara,W., Gunaratne,P., Hale,S., Hawes,A., Hernandez,J., Homsil,F., Howard,S., Huber,J., Hulyk,S., Hawes,A., Hernandez,J., Honsil,F., Howard,S., Huber,J., Hulyk,S., Hume,J., Jackson,L.E., Jackson,E., Kelly,S., Khan,U., King,L., Korvah,J., Kovar,C., Kratovic,J., Kureshi,A., Landry,N., Locier,R., Luna,R., Martinez,E., Maheshwari,M., Mapua,P., Martin,R., Martindale,A., Martinez,E., Maheshwari,M., Mapua,P., Martin,R., Martindale,A., Martinez,E., Mansey,E., Mawhiney,E., McLeed,M.P., Mawes,A., Nguyen,N., Novieto,R., Palus,F., Lucier,R., Luna,R., Mayen,A., Nguyen,N., Novieto,R., Palus,F., Martinez,B., Martinez,E., Martinez,E., Mussey,E., Mackerson,E., Nwckenkwo,S., Oghh,M., Okwnou,G., Martinez,E., Marter,R., Martinez,H., Shoshkari,M., Sisson,I., Sodergre, S., Soott,G., Shen,H., Shoshkari,M., Sisson,I., Sodergre,S., Savery,G., Sehrer,S., Mard-Moore,S., Warren,R., Washington,C., Watliag,H., Stone,H., Shoshkari,M., Sisson,R., Wang,G., Willias,M., Thomas,N., Thomas,N., Thomas,N., Thomas,N., Thomas,N., Thomas,N., Thomas,N., Martinez,E., Marter,R., Washington,C., Morley,K.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rattus norvegicus clone CH230-2A19, *** SEQUENCING IN PROGRESS ***, 76 unordered pieces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Métazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                           Submitted (17-SEP-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                               Submitted (07-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 6, 2002 this sequence version replaced gi:17940880.
                  Center project name: GACW
                                                                                                           Center: Baylor College of Medicine
                                                                                                                                    Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    * NOTE: Estimated insert size may differ from sequence length

(see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

* NOTE: This is a 'working draft' sequence. It currently

* consists of 76 contigs. The true order of the places

* is not known and their order in this sequence record is

* arbitrary. Gaps between the contigs are represented as

* runs of N, but the exact sizes of the gaps are unknown.

* This record will be updated with the finished sequence

* as soon as it is available and the accession number will

* be preserved.
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----- Summary Statistics
Sequencing vector: Plasmid;
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16984: contig
17084: gap of u
18124: contig
18224: gap of u
19324: gap of u
19344: gap of u
19444: gap of u
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contig of 1050
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JOURNAL REFERENCE AUTHORS TITLE

Worley, K.C. Direct Submission

JOURNAL

TITLE

Unpublished Direct Submission

(bases 1 to 115643)

REFERENCE

Baylor Plaza, Houston, 3 (bases 1 to 115643)

AUTHORS

Direct Submission Worley, K.C.

COMMENT

Center code: BCM

---- Genome

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JOURNAL TITLE

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VERSION
KEYWORDS
                                                          AC123419
        SOURCE
                             ACCESSION
                                           DEFINITION
                                                    COCUS
ORGANISM
                                                                                                                    Matches
                                                                                        19119
                                                                                             604 GTGNAAAAAAAAAAAAAAAAAAA 625
                                                                                      GTGNAAAAAAAAAAAAAAA 19098
                                         Rattus
Rattus norvegicus
                     AC123419
AC123419.2 GI:21902987
              HTG; HTGS_PHASE1.
      Rattus norvegicus.
                                                                                                                           Similarity
                                  us norvegicus clone CH230-248I13, 43 unordered pieces.
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AUTHORS
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             Direct Submission

Submitted (23-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
On Jul 18, 2002 this sequence version replaced gi:21240349.
                                                                                                                                                                                                                                                                                                                                                                                 Submitted (29-MAY-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                 Worley, K.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Worley, K.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (bases 1 to 126363)
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Center: Baylor College of Medicine Center code: BCM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submission
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Center project name: GXDX
Center clone name: GXDX
Center clone name: CH230-248113
Center clone name: CH230-248113
Center clone name: CH230-248113
Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap: version 0.990329
Consensus quality: 85769 bases at least Q40
Consensus quality: 89524 bases at least Q30

Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
Project Information

```
NOTE: Estimated insert size may differ from sequence length
(see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently
consists of 43 contigs. The true order of the pieces
is not known and their order in this sequence record is
arbitrary. Gaps between the contigs are represented as
runs of N, but the exact sizes of the gaps are unknown.

This record will be updated with the finished sequence
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            requests: clonerequestesanger.ac.uk

On Apr 22, 2001 this sequence version replaced gi:12329466.

During sequence assembly data is compared from overlapping clones.

Where differences are found these are annotated as variations
together with a note of the overlapping clone name. Note that the
variation annotation may not be found in the sequence submission
corresponding to the overlapping clone, as we submit sequences with
only a small overlap as described above.

This sequence was finished as follows unless otherwise noted: all
regions were either double-stranded or sequenced with an alternate
chemistry or covered by high quality data (i.e., phred quality >=
30); an attempt was made to resolve all sequencing problems, such
as compressions and repeats; all regions were covered by at least
one plasmid subclone or more than one M13 subclone; and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 22; Conserv
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                             Submitted (20-APR-2001) Sanger Centre, Hinxton, Cambridgeshire CB10 1SA, UK. E-mail enquiries: humquery@sanger.ac.uk Clone
                                                                                                                                                                                                                                                                                                                                AL445433 138918 bp DNA linear PRI 20-APR-2001
Human DNA sequence from clone RP11-234N17 on chromosome 1, complete
                                                                                                                                                                                                     Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      http://www.sanger.ac.uk/HGP/Chrl
RP11-234N17 is from the library RPCI-11.1 constructed by the group
of Pieter de Jong. For further details see
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                http://www.sanger.ac.uk/Projects/C_elegans/wormpep This sequence was generated from part of bacterial clone contigs of human chromosome 1, constructed by the Sanger Centre Chromosome 1 Mapping Group. Further information can be found at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence. The true right end of clone RP5-846Fl3 is at 100 in this
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 abbreviations are used to associate primary accession numbers given in the feature table with their source databases: Em:, EMBL; Sw:,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         http://www.chori.org/bacpac/home.htm
VECTOR: pBACe3.6
                            /note-match: GSS: Em:B98754 15733. 16309
                                                                                                                                                                                                  /note-"MIR repeat: matches 37.
11749. .12081
                                                                                                           /note="MER5B repeat: matches 5. .178 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="match: GSS: Em:AQ042814"
complement(3711. .3794)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      complement(3711
                                                                                                                                                           /note="L1ME1 repeat: matches 5764. .6118 of consensus"
                                                                                                                                                                                                                                                               /note="MLT1B repeat: matches 1. .390 of consensus"
                                                                                                                                                                                                                                                                                                              /note="AluJo repeat: matches 1. .310 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                       'note-"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="match: GSS: Em:AQ519708" complement(3700. .3925)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="MIR repeat: matches 79.
complement(3065, .3515)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="Sequence confirmed by AC020693 and AC022823 both
sequenced by WIBR."
                                                                                                                                                                                                                                                                                                                                                                             'note="
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'note-"MIR repeat: matches 8. .166 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           note="match: GSS: Em:AZ694502"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             note="match: GSS: Em:AQ078674"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note-"match: GSS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
/chromosome="1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note-"MIR repeat: matches 19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'note="CpG island"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   clone_lib="RPCI-11.1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'clone="RP11-234N17"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .138918
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L2 repeat: matches 2127. .2710 of consensus" .17208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .5739
                                                                                                                                                                                                                                                                                                                                                                     AluSq repeat: matches 1. .292 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                       AluJo repeat: matches 1. .286 of consensus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AIR repeat: matches 81. .145 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 repeat: matches 2650. .2749 of consensus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .3925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Em: B38703"
                                                                                                                                                                                                                  .251 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .202 of
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                                                                                                                                                                                                                                                                                                                                                                            /note="MIR repeat: matches 12. .151 of consensus" 28154. .28513
/note="L1MA9 repeat: matches 6037. .6308 of consensus" 34110. .34365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22808 . .22900
/note="MIR repeat: matches 48 . .142
/note=ment(22989 . .23434)
complement(22989 . .23434)
                                                                                                             consensus
                                                                                                                        /note="THE1B-INTERNAL repeat: matches 1.
                                                                                                                                                                                                                                                                                                                                                                                                                     27905. .28044
                                               /note="THE1B repeat: matches 1. .364 of consensus" 33482. .33632
                                                                                                                                                                    /note="THE1B repeat: matches 1. .364 of consensus"
                                                                                                                                                                                                                                                            /note="L1 repeat: matches 3750. .4361 of consensus"
                                                                                                                                                                                                                                                                                                        /note≖"LlM3a repeat: matches 913. .1541 of consensus"
                                                                                                                                                                                                                                                                                                                                              /note="LlM3a repeat: matches 549. .946 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27801.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="L1MA8 repeat: matches 6006. .6261 of consensus"
27418. .27800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="Alur repeat: matches 81. .306 of consensus"
27174. .27417
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="LlMA8 repeat: matches 6261. .6287 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="MLTIF repeat: matches 359. .568 of consensus" complement(22703. .23209)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="L1MC4 repeat: matches 5254, .6928 of consensus"
20828, .20885
/note="29 copies 2 mer tt 70% conserved"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="AluJo repeat: matches 1..162 of consensus" 18259. .18675
/note="MSTB repeat: matches 6..425 of consensus" 18708. .19182
                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="11MA8 repeat: matches 5924. .6006 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26949
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   note-"MSTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="L2 repeat: matches 1704. .2180 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="L2 repeat: matches 1176. .1564 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="14 copies 2 mer gt 96% conserved"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note="L2 repeat: matches 2532.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note="match: GSS: Em:AQ758666"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'note="AluJb repeat:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "U5 repeat: matches 1. .46 of consensus" .25839
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                repeat: matches 1. .424 of consensus"
                                                                                                                                                                                                                 repeat: matches 4704. .6037 of consensus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       repeat: matches 1. .462 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    repeat: matches 1. .111 of consensus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         repeat: matches 5011. .5403 of consensus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .2750 of consensus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .6142 of consensus"
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             Muzny, D. Marie . Metzker, M. Lee., Abramzon, S., Adams, C., Alder, J., Allen, C., Allen, H., Alsbrooks, S., Amin, A., Angulano, D., Angulano, D., Angulabechi, V., Aoyagi, A., Ayodeji, M., Baca, E., Baden, H., Balakin, D., Bandarannaike, D., Barber, M., Barnstead, M., Benahmed, F., Baldwin, D., Bandarannaike, D., Barber, M., Barnstead, M., Benahmed, F., Biswalo, K., Blair, J., Blankenburg, K., Blyth, P., Brown, M., Biswalo, K., Blair, J., Blankenburg, K., Blyth, P., Brown, M., Biswalo, K., Blair, J., Blankenburg, K., Blyth, P., Brown, M., Biswalo, K., Bulay, C., Burch, P., Burrell, K., Calderon, E., Cardenas, V., Carter, K., Cavazos, I., Ceassar, H., Center, A., Chen, C., Burch, P., Burchl, P., Cox, C., Coyle, M., Cree, A., D'Souza, L., Chacko, J., Chen, C., Coyle, M., Cree, A., D'Souza, L., Chacko, J., Cox, C., Coyle, M., Cree, A., D'Souza, L., Davils, C., Davils, M., Lovis, C., Davy, Carroll, L., De Anda, C., Dederich, D., Davils, M., Loyan, S., Durn, A., Durbin, K., Duval, B., Eaves, K., Delgando, O., Denson, S., Deramo, C., Ding, Y., Dinh, H., Divya, K., Deger, H., Durya, K., Durah, A., Escotto, M., Eugene, C., Evans, C.A., Falls, T., Fan, G., Egan, A., Escotto, M., Eugene, C., Evans, C.A., Falls, T., Fan, G., Egan, A., Escotto, M., Ganta, R., Garcia, A., Gerrer, M., Gebregeorgis, E., Geet, K., Gill, R., Garcia, A., Gerrer, M., Gebregeorgis, E., Geet, K., Gill, R., Garcia, A., Gerrer, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra, W., Guerra
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Rattus norvegicus clone CH230-19217, *** SEQUENCING IN PROGRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
Lorensuhewa,L., Loulseged,H., Lozado,R.J., Lu,X.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (bases 1 to 158728)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="MIR repeat: matches 3. .260 of consensus" 34736. .38491
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38655. .4018
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42450. .42781
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42418 . .42449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="L1PA8 repeat: matches 5942. .6163 of consensus"
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42139. .42417
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Mangum, B., Mapua, P., Martin, K., Martin, R., Martinez, E.,
Mangum, B., Mapua, P., Martin, K., Martin, R., Martinez, E.,
Manviney, S., Mcleod, M., Mcneill, T., Meenen, E., Milosavljevic, A.,
Miner, G., Minja, E., Montemayor, J., Moore, S., Morgan, M., Morris, K.,
Miner, S., Munidasa, M., Murphy, M., Nair, L., Nankervis, C., Neal, D.,
Morris, S., Munidasa, M., Murphy, M., Nair, L., Nankervis, C., Neal, D.,
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Popovic, D., Primus, E., Pu, L.-L., Puazo, M., Quiroz, J., Rachlin, E.,
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Rose, R., Ruiz, S.J., Sanders, W., Savery, G., Scherer, S.,
Sotott, G., Shatsman, S., Shen, H., Shetry, J., Shvartsbeyn, A.,
Sisson, I., Sitter, C.D., Smajs, D., Sneed, A., Sodergren, E.,
Song, X.-Z., Sorelle, R., Sosa, J., Steinle, M., Strong, R., Sutton, A.,
Song, X.-Z., Sorelle, R., Sosa, J., Steinle, M., Strong, R., Sutton, A.,
Svatek, A., Tabor, P., Taylor, C., Taylor, T., Thomas, N., Thomas, N.,
Thomas, N., Wailliams, G., Wallson, R., Wieczyk, R.,
Warren, R., Wei, X., White, F., Williams, G., Willson, R., Wieczyk, R.,
Weoden, H., Worley, K., Wright, D., Wright, R., Wu, J., Yakub, S.,
Yen, J., Yoon, L., Yoon, V., Yu, F., Zhang, J., Zhou, J., Zhou, X.,
Palao, S., Dunn, D., von Niederhausern, A., Weiss, R., Smith, D.R.,
Palao, S., Smith, D., Weinstock, G. and Gibbs, R.A.
                                     Worley,K.C.
Direct Submission
Submitted (07-MAR-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Maheshwari,M., Mahindartne,M., Mahmoud,M., Malloy,K., Mangum,A.,
                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished
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(bases 1 to 158728)
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JOURNAL AUTHORS

Submitted (24-Aug-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA on Aug 14, 2002 this sequence version replaced gi:21731615. Direct Submission Rat Genome Sequencing Consortium.

-- Genome Center

Center: Baylor College of Medicine Center code: BCM

Center clone name: CH230-19217

Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 92840 bases at least Q40
Consensus quality: 100722 bases at least Q20
Consensus quality: 105423 bases at least Q20

\* NOTE: Estimated insert size may differ from sequence length

\* (see http://www.hgsc.bcm.tmc.edu/docs/Genbank\_draft\_data.html).

\* NOTE: This is a 'working draft' sequence. It currently

\* consists of 80 contigs. The true order of the places

\* is not known and their order in this sequence record is

\* arbitrary. Gaps between the contigs are represented as

\* runs of N, but the exact sizes of the gaps are unknown.

\* This record will be updated with the finished sequence

\* This record will be updated with the finished sequence

\* as soon as it is available and the accession number will

\* be preserved.

1219 1319 2615 2715 2715 3715 3815 5074 5174 1218: contig of 1218 bp in length
1318: gap of unknown length
2614: contig of 1296 bp in length
2714: gap of unknown length
3714: contig of 1000 bp in length
3814: gap of unknown length
5073: contig of 1259 bp in length
5173: gap of unknown length
6512: contig of 1339 bp in length

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20817: gap of
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36041:
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JOURNAL REFERENCE
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Birren, B., Linton, L., Nusbaum, C. and Lander, E. Homo sapiens chromosome 1, clone RP11-76A15
                       Submitted (06-FEB-2000) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA 1 (bases 1 to 160008)

Birren,B., Linton,L., Nusbaum,C., Lander,E., Abraham,H., Allen,N., Anderson,S., Baldwin,J., Barna,N., Bastien,V., Beda,F., Boguslavkiy,L., Boukhgalter,B., Brown,A., Burkett,G., Campopiano,A., Castle,A., Choepel,Y., Colangelo,M., Collins,S., Collymore,A., Cooke,P., DeArellano,K., Dewar,K., Diaz,J.S., Dodge,S., Domino,M., Doyle,M., Ferreira,P., FitzHugh,W., Gage,D., Galagan,J., Gardyna,S., Ginde,S., Goyette,M., Graham,L.,
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Galagan,J., Gardyna,S., Ginde,S., Goyette,M., Graham,L.,
Grand-Pierre,N., Grant,G., Hagos,B., Heaford,A., Horton,L.,
                                                                                                                                                                                                                                                                                                                 Direct Submission
                                                                                                                                                                                                                                                                                                                                         Zimmer, A and Zody, M.
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Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Roy,A., Santos,R., Schauer,S., Severy,F., Crunce, Talamas,J., Stange-Thomann,N., Stojanovic,N., Subramanian,A., Talamas,J., Tesfaye,S., Theodore,J., Tirrell,A., Travers,M., Trigilio,J., Tesfaye,S., Theodore,J., Wilson,B., Wu,X., Wyman,D., Ye,W.J., Vassillev,H., Viel,R., Vo,A., Wilson,B., Wu,X., Wyman,D., Ye,W.J., Young,G., Zainoun,J., Zimmer,A. and Zody,M.
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http://ftp.genome.washington.edu/RM/RepeatMasker.html
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Klein,J., LaRocque,K., Lamazares,R., Landers,T., Lehoczky,J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      as soon as it is available and the accession number will be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown This record will be updated with the finished sequence
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1213 2428:
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2 9441: σ=
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2428: contig of 1216
2528: gap of 100 bp
3940: contig of
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15511: con
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511: gap of 100 bn
17718: conti
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20410: contig of 2592 bp in length
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5729:
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120993 131870: cont
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59083 63451: contig of 4369 bp in length
63452 63551: gap of 100 bp
63552 68086: contig of 4535 bp in length
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131971 142739: cont
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31661 35351: contig of 3691 bp in
35352 35451: gap of 100 bp
35452 38476: contig of 3025 bp in
38477 38576: gap of 100 bp
38577 41099: contig of 2523 bp in
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5830. .7163
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4 58982:
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14758: contig of 3468 bp in length
14758: gap of 100 bp

51427: contig of 3869 bp in length
51527: gap of 100 bp
54783: contig of 3256 hr
1483: gap of 3256 hr
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/note="assembly_fragment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24651:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23006:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100 bp
100 bp
100 bp
19082; gap of
63451; con++
1511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12839: gap of 
160008: cont
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26805: contig of 2154 bp in length

905: gap of 100 bp

29287: contig of 2382 bp in length

87: gap of 100 bp

31560: contig of 2173 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24551:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98963:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16: gap of 100 bp
2020: contig of 3834 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'assembly_fragment"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gap of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gap of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gap of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ap or 100 bp
contig of 3974 bp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             p of 100 bp
contig of 7320 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              contig of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              contig of 17169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ap of 100 bp
contig of 10769 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                contig of 10878 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100 bp
104 bp in 7
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f 2523 bp in length
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RESULT 40
AC020693
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            VERSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  397 GAAAAACACAGCATGAGAACAC 418
RS Birren, B. Linton, L. Nusbaum, C., Lander, E., Abraham, H., Allen, N., Anderson, S., Baldwin, J., Barna, N., Bastien, V., Beda, F., Anderson, S., Baldwin, J., Barna, N., Bastien, V., Beda, F., Boukhgalter, B., Brown, A., Burkett, G., Callarkiy, L., Boukhgalter, B., Brown, A., Burkett, G., Collins, S., Campopiano, A., Castle, A., Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Domino, M., Doyle, M., Ferreira, P., Fitzhugh, W., Gage, D., Galagan, J., Gardyna, S., Ginde, S., Goyette, M., Graham, L., Galagan, J., Gardyna, S., Ginde, S., Goyette, M., Graham, L., Galagan, J., Gardyna, S., Ginde, S., Goyette, M., Graham, L., Howland, J.C., Iliev, I., Johnson, R., Jones, C., Kann, L., Karatas, A., Kieln, J., Lancoque, K., Lanazares, R., Landers, T., Lehoczky, J., Levine, R., Lieu, C., Liu, G., Locke, K., Macdonald, P., Marquis, N., McCarthy, M., McEwan, P., McGurk, A., McKennan, K., McPheeters, R., Meldrim, J., Weneus, L., Mihova, T., Miranda, C., Mlenga, V., Morrow, J., Murphy, T., Naylor, J., Norman, C.H., O'Connor, T., O'Donnell, P., O'Neil, D., Olivar, T. M., Oliver, J., Peterson, K., Pierre, N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anderson, S., Baldwin, J., Barna, N., Beckerly, R., Beda, F.,
BoguslavKiy, L., Boukhgalter, B., Brown, A., Burkett, G., Castle, A.,
Choepel, Y., Colangelo, M., Collins, S., Collymore, A., Cooke, P.,
Pearellano, K., Dewar, K., Domino, M., Doyle, M., Fenestor, J.,
Frizhugh, W., Forrest, C., Gage, D., Galagan, J.,
Gardyna, S., Grant, G., Hagos, B., Heaford, A., Horton, L.,
Gardyna, S., Grant, G., Hagos, B., Heaford, A., Horton, L.,
Howland, J.C., Johnson, R., Jones, C., Kann, L., Karatas, A., Klein, J.,
Howland, J.C., Johnson, R., Jones, C., Kann, L., Karatas, A., Klein, J.,
Howland, J.C., Johnson, R., Jones, C., Kann, L., Karatas, A., Klein, J.,
Howland, J.C., Johnson, R., Jones, C., Kann, L., Karatas, A., Klein, J.,
Howland, J.C., Johnson, R., Jones, C., Kann, L., Karatas, A., Klein, J.,
Macdonald, P., Marquis, N., McEwan, P., McGurk, A., McKernan, K.,
McDheeters, R., Meldrim, J., Meneus, L., Morrow, J., Naylor, J.,
Morman, C.H., O'Connor, T., O'Donnell, P., Olivar, T.M., Peterson, K.,
Pierre, N., Pisani, C., Pollara, V., Raymond, C., Riley, R., Rothman, D.,
Rova, A., Santone R., Severy D., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., Santone R., San
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
Submitted (08-JAN-2000) Whitehead Institute/MIT Center for Genome Sesearch, 320 Charles Street, Cambridge, MA 02141, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ROY,A., Santos,R., Severy,P., Spencer,B., Stange-Thomann,N., Stojanovic,N., Subramanian,A., Talamas,J., Tesfaye,S., Theodore,J. Tirrell,A., Vassiliev,H., Viel,R., Vo,A., Wu,X., Wyman,D., Ye,W.J. Zimmer,A. and Zody,M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 163932)
Birren,B., Linton,L., Nusbaum,C. and Lander,E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Birren, B., Linton, L., Nusbaum, C., Lander, E., Abraham, H., Allen, N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACO20693.3 GI:7249430
HTG; HTGS_PHASE1; HTGS_DRAFT.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACO20693 163932 bp DNA
Homo sapiens chromosome 6 clone RP11-574H19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AC020693
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15612. .17718
/note="assembly_fragment"
17819. .20410
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9442. .10979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="assembly_fragment"
11080. .12887
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9 map 6, WORKING DRAFT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Theodore, J.,
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JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (24-AUC-2002) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA on Mar 16, 2000 this sequence version replaced gi:6730833.

All repeats were identified using RepeatMasker:
Smit, A.F.A. & Green, P. (1996-1997)
http://ftp.genome.washington.edu/RM/RepeatMasker.html
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 2756: contig of 2756 bp in length 2757 2856: gap of 100 bp 2857 7124: contig of 4268 bp in length 7125 7224: gap of 100 bp 1725 14958: contig of 734 bp in length 14959 15058: gap of 100 bp 15059 21054: contig of 5996 bp in length 21055 21154: gap of 100 bp 1520: contig of 100 bp 1520: contig of 100 bp 1520: contig of 100 bp 158638: gap of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100 bp 158638: contig of 100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequencing vector: M13; M77815; 100% of reads Chemistry: Dye-terminator Big Dye; 100% of reads Assembly program: Phrap; version 0.960731 Consensus quality: 159231 bases at least Q40 Consensus quality: 161639 bases at least Q20 Consensus quality: 162468 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        58638 58737: gap of 100 bp
58738 110374: contig of 51637 b
110375 110474: gap of 100 bp
110475 163932: contig of 53458 k
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Insert size: 16323; sum-of-contigs
Quality coverage: 5.4 in Q20 bases; agarose-fp
Quality coverage: 5.2 in Q20 bases; sum-of-contigs
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Center clone name: 574_H_19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: sequence_submissions@genome.w1.mit.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Web site: http://www-seq.wi.mit.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Center code: WIBR
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                                                                                                7225. .14958
                                                                                                                                                                                                                                                vector_
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="RP11-574H19"
/clone_lib="RPCI-11 Human Male BAC"
                                     /note="assembly_fragment"
                                                                                                                                           /note="assembly_fragment"
                                                                                                                                                                                                                                                                                                                                                      note-"assembly_fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /chromosome="6"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /db_xref="taxon:9606"
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110374: contig of 51637 bp in length
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.21054
                                                                                                                                                                                                                                                side:right"
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RS MuznyD.M., Adams,C., Adio-Oduola,B., Ali-osman,F.R., Allen,C., Alsbrooks,S.L., Amaratunde,H.C., Are,J.R., Ayele,M., Banks,T., Blabaria,J., Benton,J., Blamage,K., Blankenburg,K., Bonnin,D., Bouck,J., Bowle,S., Brieva,M., Brown,E., Brown,M., Bryant,N.P., Bouck,J., Burch,P., Burkett,C., Burrell,K.L., Byrd,N.C., Carron,T.F., Carter,M., Cavazos,S.R., Chacko,J., Chavez,D., Chen,G., Chen,R., Chen,z., Chowdhry,I., Christopoulos,C., Chen,G., Chen,R., Chen,Z., Coyle,M.D., Dathorne,S.R., David,R., Davila,M.L., Davis,C., Coyle,M.D., Dathorne,S.R., David,R., Davila,M.L., Davis,C., Davy-Carroll,L., Dederich,D.A., Delaney,K.R., Delgado,O., Denn,A.L., Ding,Y., Dinh,H.H., Delaney,K.R., Delgado,O., Denn,A.L., Ding,Y., Dinh,H.H., Delaney,K.R., Delgado,O., Denn,A.L., Ding,Y., Dinh,H.H., Falls,T., Ferraguto,D., Flaggy,N., Ford,J., Foster,P., Frantz,P., Falls,T., Gao,J., Garcia,A., Garner,T., Garza,N., Gill,R., Gabisi,A., Gao,J., Garcia,A., Garner,T., Garza,N., Gill,R., Garris,C., Harris,C., Harris,K., Hart,M., Havlak,P., Hawes,A., Hernandez,J., Homsi,F., Howard,S., Huber,J., Hulk,S., Hune,J., Jackson,L.E., Janobson, B., 11a,Y., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon, B., Tohnon
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Mammalia; Eutheria; Rodentia; Sclurognathi; Muridae; Murinae;
Sutton,A., Svatek,A., Tabor,P., Tamerisa,A., Tamerisa,K., Tang,H., Tansey,J., Taylor,C., Taylor,T., Telfrod,B., Thomas,N., Thomas,S., Usmani,K., Vasquez,L., Vera,V., Villalon,D., Vinson,R., Wang,Q.,
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110475. .163932
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31621. .58637
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Submitted (17-SEP-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plazz, Houston, TX 77030, USA
3 (bases 1 to 194402)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted (11-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Baylor Plaza,
On Jul 10, 200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Worley, K.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NOTE: Estimated insert size may differ from sequence length (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

NOTE: This is a 'working draft' sequence. It currently consists of 77 contigs. The true order of the pieces is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              as soon as it is available and the accession number will be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jul 10, 2002 this sequence version replaced gi:17943838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap; version 0.990329
Consensus quality: 127428 bases at Least Q40
Consensus quality: 136154 bases at least Q30
Consensus quality: 142855 bases at least Q30
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Center clone name: CH230-10H22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Center code: BCM
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ACCESSION VERSION

DEFINITION AC096152/c

Rattus

KEYWORDS

SOURCE

Rattus norvegicus Norway rat HTG; HTGS\_PHASE1. AC096152.4 GI:21723293

REFERENCE

Rattus

(bases 1 to 194402)

AUTHORS

BASE COUNT ORIGIN

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vector\_side:left" 58738. .110374

clone\_end:SP6

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misc\_feature

Matches

Conservative

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Query Match Best Local Similarity

Db 124291 GAAAAACACAGCATGAGAACAC 124312

397 GAAAAACACAGCATGAGAACAC 418

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Wang,S., Ward-Moore,S., Warren,R., Washington,C., Watlington,S., Williams,G., Williamson,A., Wleczyk,R., Wooden,S., Worley,K., Wu,C., Wu,Y., Wu,Y.F., Zhou,J., Zorrilla,S., Nelson,D., Weinstock,G. and Gibbs,R.
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of 1035

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                                                                                                         Alabrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbaria, J., Benton, J., Binage, K., Blankenburg, K., Bonks, T., Barbaria, J., Benton, J., Binage, K., Blankenburg, K., Bonks, T., Burkett, C., Burch, H., Bryant, N.P., Burkett, C., Burch, E., Brown, M., Bryant, N.P., Bunks, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Bunks, J., Cheve, J., Chacke, J., Chacke, J., Chavez, D., Chen, Z., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Chen, J., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwalte, K.J., Drayer, H., Dugan-Rocha, S., Durbin, K.J., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Gorrell, J.H., Guevara, W., Gunaratne, P., Hale, S., Hamilton, K., Harris, K., Harris, K., Hart, M., Haylak, P., Hawes, A., Hernandez, J., Hodgson, A., Hogues, M., Hollowy, C., Hellins, B., Hensi, F., Howard, S., Lucier, R., Julyk, S., Hume, J., Jackson, L.E., Homsi, F., Howard, S., Kanlyu, K., Kovah, J., Kovar, C., Kratovic, J., Kureshi, A., Landry, N., Leal, B., Lewis, L.C., Levis, L., Lichtarge, O., Lieu, C., Liu, J., Liu, W., Loulseed, H., Lozado, R.J., Lu, X., Lucier, R., Luna, R., Martinez, E., Mansen, J., Wasten, R., Waster, A., Wartinez, E., Mansen, C., Miner, Z., Mitchell, T., Mohabbat, K., Morgan, M., Morris, S., Massey, E., Machiney, E., McLeod, M.P., Meadon, M., Nguyen, A., Nguyen, N., Nickerson, E., Necken, N., Pace, A., Payton, B., Peery, J., Perez, L., Filmer, J., Pulles, M., Ren, Y., Rives, M., Royles, A., Roylas, A., Roylubkan, I., Rolfe, M., Ru, K., Scherer, S., Scott, G., Shen, H., Shooshtari, N., Sisson, I., Sodergen, S., Seatke, A., Tabor, P., Tamerisa, A., Tamerisa, A., Tang, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, S., Worley, K., Wardon, S., Worley, K., Wardon, S., Worley, K., Wardon,
Unpublished 2 (bases 1
                                                   Direct Submission
                                                                          Wu,C., Wu,Y., Wu,Y.F., Zhou,J., Zorrilla,S., Nelson,D., Weinstock,G. and Gibbs,R.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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98378: gap of u
101131: contig
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Submitted (13-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (23-JAN-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA 3 (bases 1 to 196863)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Worley, K.C.
Direct Submission
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Center clone name: CH230-101C21
Center clone name: CH230-101C21
Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap; version 0 990329
Consensus quality: 153108 bases at least Q40
Consensus quality: 159647 bases at least Q20
Consensus quality: 166406 bases at least Q20
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On Jul 11, 2002
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NOTE: This is a 'working draft' sequence. It currently consists of 62 contigs. The true order of the pieces is not known and their order in this sequence record is
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Birren, B., Nusbaum, C., Lander, E., Ali, A., Allen, N., Anderson, S., Barna, N., Bastien, V., Bloom, T., Boguslavkiy, L., Boukhgalter, B., Camarata, J., Chang, J., Chazaro, B., Choepel, Y., Collymore, A., Cook, A., Cooke, P., DeArellano, K., Dewar, K., Diaz, J.S., Dodge, S., Faro, S., Ferreira, P., FitzGerald, M., Gage, D., Galagan, J., Gardyna, S., Gord, S., Graham, L., Grand-Pierre, N., Hagos, B., Horton, L., Hulme, W., Iliev, I., Johnson, R., Jones, C., Kamat, A., Katatas, A., Kells, C., Landers, T., Levine, R., Lindblad-Toh, K., Liu, G., Macchan, C., Macdonald, P., Major, J., Matthews, C., McCarthy, M., Meldrim, J., Menus, L., Mihova, T., Mlenga, V., Marchy, M., Meldrim, J., Menus, L., Mihova, T., Mlenga, V., Marchy, M., Meldrim, J., Menus, L., Mihova, T., Mlenga, V., Marchy, M., Meldrim, J., Menus, L., Minova, T., Mienga, V., Marchy, M., Marthey, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan, C., Macchan
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                                    Submitted (21-AUG-2002) Whitehead Institute/MIT Center for Genome
Research, 320 Charles Street, Cambridge, MA 02141, USA
                                                                                                                          Direct Submission
                                                                                                                                                                                                  Zembek, L., Zimmer, A. and Zody, M.
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Birren,B., Nusbaum,C. and Lander,E.
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On Aug 21, 2002 this sequence version replaced gi:17061674. All repeats were identified using RepeatMasker: Smit, A.F.A. & Green, P. (1995-1997) http://ftp.genome.washington.edu/RM/RepeatMasker.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  * arbitrary. Gaps between the contigs are represented as 
* runs of N, but the exact sizes of the gaps are unknown.
* This record will be updated with the finished sequence
* as soon as it is available and the accession number will
* be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              * NOTE: This is a 'working draft' sequence. It currently * consists of 25 contigs. The true order of the pieces * is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      * 8314: contig of 1772 bp in length

* 8315 8414: gap of 100 bp

8415 10954: contig of 2540 bp in length

10955 11054: gap of 100 bp

11055 12504: contig of 3550 bp in length

14505 14604: gap of 100 bp

14505 14604: gap of 100 bp

14605 19881: contig of 5277 bp in length

19882 19881: gap of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequencing vector: Plasmid; n/a; 100% of reads Chemistry: Dye-terminator Big Dye; 100% of reads Assembly program: Phrap; version 0.960731 consensus quality: 191319 bases at least Q40 consensus quality: 194846 bases at least Q30 consensus quality: 195925 bases at least Q20 consensus quality: 195925 bases at least Q20 Insert size: 198000; agarose-fp Insert size: 1980443; sum-of contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Quality coverage: 5.8 in Q20 bases; agarose-fp Quality coverage: 5.8 in Q20 bases; sum-of-contigs
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Center clone name: 334_D_14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Center code: WIBR
                                                                                                                                                                                                                              63373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ----- Summary Statistics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Center: Whitehead Institute/ MIT Center for Genome Research
                                                               99214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    142 761: contig of 620 bp in length
762 861: gap of 100 bp
862 1803: contig of 942 bp in length
1804 1903: gap of 100 bp
1904 3084: contig of 1181 bp in length
3085 3184: gap of 100 bp
3185 3184: gap of 100 bp
                                                                                                                                                                                                                                                                                                                                          44875
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4761 6
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4661 4760: gap of 100 bp
4761 6442: contig of 1682 bp in length
6443 6542: gap of 100 bp
  135934: gap of
151141: cont
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135834: contig of 36621 bp in length
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                                                                                                                                                                                                                256: gap of 100 bp
63272: contig of 10016 bp in length
372: gap of 100 bp
73192: contig of 9820 bp in length
                                                                                                           106: gap of 100 bp
19113: contig of 13007 bp in length
                                                                                                                                                               192: gap of 100 bp 86006: contig of 12714 bp
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53156: contig of 8282 bp in length
                                                                                                                                                                                                                                                                                                                                                                                        )51: gap of 100 bp
44774: contig of 7723 bp in length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    81: gap of 100 bp 24662: contig of 4681 bp in length
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ap of 100 bp contig of 15207 bp in length
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Db 122888 AGAACACAGTAAATGAATAAAA 122867
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151242 162188: contig of 10947 bp in length
162189 162288: gap of 100 bp
162289 175380: contig of 14092 bp in length
176381 176480: gap of 100 bp
176481 187921: contig of 11441 bp in length
187922 188021: gap of 100 bp
188022 198843: contig of 10822 bp in length.
Location/Qualifiers
                                                                  Conservative
                                                                                                                                                                                                                                                                     /note="assembly_fragment"
135935, .151141
                                                                                                                                                                                                                                                                                                     /note-"assembly_fragment"
99214. .135834
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63373. .73192
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53257. 63272
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/db_xref="taxon:10090"
/clone="RP23-334D14"
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Pred. No. 0.52;
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                                                                                                                                  TITLE
 Submitted (29-MAY-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA 3 (bases 1 to 206648)
                                                                    Direct Submission
                                                                                 Worley, K.C.
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Bryant, N., Buhay, C., Burch, P., Burrell, K., Caideron, E.,
Cardenas, V., Carter, K., Cavazos, I., Ceasar, H., Center, A.,
Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chen, Z.,
Cleveland, C., Cockrell, R., Cox, C., Coyle, M., Cree, A., D'Souza, I.,
Cleveland, C., Cockrell, R., Cox, C., Coyle, M., Cree, A., D'Souza, I.,
Davila, M.L., Davis, C., Deramo, C., Ding, Y., Dinh, H., Divya, K.,
Delgado, O., Denson, S., Deramo, C., Ding, Y., Dinh, H., Divya, K.,
Delgado, O., Denson, S., Deramo, C., Ding, Y., Dinh, H., Divya, K.,
Deraper, H., Dugan-Rocha, S., Dunn, A., Durbin, K., Daval, B., Eaves, K.,
Egan, A., Escotto, M., Eugene, C., Evans, C.A., Falls, T., Fan, G.,
Fernandez, S., Finley, M., Flagg, N., Forbes, L., Foster, M., Foster, P.,
Fraser, C.M., Gabisi, A., Ganta, R., Garcia, A., Garrer, T., Garza, M.,
Gebregeorgis, E., Geer, K., Gill, R., Grady, M., Guerra, W., Guevara, W.,
Harvey, Y., Havlak, P., Hawes, A., Henderson, N., Hernandez, J.,
Hernandez, R., Hines, S., Hladun, S.L., Hodgson, A., Hogues, M.,
Hellins, B., Howells, S., Hulyk, S., Hume, J., Idlebird, D., Jackson, A.,
Jackson, L., Jacob, L., Henderson, N., Hernandez, J.,
Hernandez, R., Kally, S., Kally, S., Khan, Z., King, L., Kovar, C.,
Kowls, C., Kraft, C.L., Lebow, H., Levan, J., Idlebird, D., Jackson, A.,
Jackson, L., Jacob, L., Jang, H., Johnson, B., Johnson, R., Jolivet, A.,
Karpathy, S., Kelly, S., Kelly, S., Khan, Z., King, L., Kovar, C.,
Kowls, C., Kraft, C.L., Lebow, H., Levan, J., Lewis, L., Li, Z., Liu, J.,
Liu, J., Liu, W., Liu, Y., London, P., Longacre, S., Morgan, M., Morris, K.,
Mangum, B., Mapun, P., Martin, K., Martin, R., Martin, E.,
Manum, B., Mapun, P., Martin, K., Martin, K., Martin, K.,
Martin, S., Munidas, M., Norris, S., Norgan, M., Morris, K.,
Morris, S., Munidas, M., Norris, S., Nwaren, S., Norgan, M., Norris, S., Nachenen, E., Pull, H., Natry, M., Ren, Y.,
Reeves, K., Rejter, M., Reigh, R., Setty, S., Scherer, S.,
Song, X., Scherer, S., Shen, H., Shetty, J., Sherris, S.,
Marten, S., Shen, H., Shetty, J., Sharrisbeyn,
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Biswalo,K., Blair,J., Blankenburg,K., Blyth,P., Brown,M.,
Bryant,N., Buhay,C., Burch,P., Burrell,K., Calderon,E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Muzny,D.Marie., Metzker,M.Lee., Abramzon,S., Adams,C., Al
Allen,C., Allen,H., Alsbrooks,S., Amin,A., Angulano,D.,
Anyalebechi,V., Aoyagi,A., Ayodeji,M., Baca,E., Baden,H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rattus norvegicus
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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*** SEQUENCING IN PROGRESS ***,
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(see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).

* NOTE: This is a 'working draft' sequence. It currently

* consists of 94 contigs. The true order of the pieces

is not known and their order in this sequence record is

* arbitrary. Gaps between the contigs are represented as

* runs of N, but the exact sizes of the gaps are unknown.

This record will be updated with the finished sequence

* as soon as it is available and the accession number will

* be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (24-AUG-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 2, 2002 this sequence version replaced gi:21239864.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rat Genome Sequencing Consortium. Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Summary Statistics
Sequencing vector: Plasmid,
Chemistry: Dye-terminator Big Dye: 100% of re
Assembly program: Phrap; version 0.990329
Consensus quality: 135401 bases at least Q40
Consensus quality: 143073 bases at least Q30
Consensus quality: 149355 bases at least Q20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Center project Information Center project name: GLJV Center clone name: CH230-68C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Web site: http://www.hgsc.bcm.tmc.edu/
Contact: hgsc-help@bcm.tmc.edu
------ Project Information
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                                                                                                                                                                                                                                                                                  Submitted (10-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA on Jul 9, 2002 this sequence version replaced gi:17941347.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Direct Submission
Submitted (15-SEP-2001) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
3 (bases 1 to 209095)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rattus norvegicus
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AC094585.3 GI:21716318
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished
2 (bases 1
                                                                                                                                                                                                                                                                                                                                                                                                                                          Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (bases 1 to 209095)
                     Center clone name: CH230-4B10
                                                                                                                              Web site: http://www.hgsc.bcm.tmc.edu/Contact: hgsc-help@bcm.tmc.edu
                                                                                                                                                                                                 Center: Baylor College of Medicine Center code: BCM
                                                                                                                                                                                                                                                                        --- Genome Center
Summary Statistics
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Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 167289 bases at least 040
Consensus quality: 178589 bases at least 030
Consensus quality: 178689 bases at least 020
Consensus quality: 178689 bases at least 020

**NOTE: Estimated insert size may differ from sequence length
**(see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).*

**NOTE: This is a 'working draft' sequence. It currently
**consists of 50 contigs. The true order of the pieces
** is not known and their order in this sequence record is
** arbitrary. Gaps between the contigs are represented as
** runs of N, but the exact sizes of the gaps are unknown.
** This record will be updated with the finished sequence
** as soon as it is available and the accession number will
** be preserved.

** 1113
1212: gap of unknown length
** 2337
2336: gap of unknown length
** 2337
3350: contig of 1244 bp in length
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8841 8941 10776 10876 11982 12022 12077 13778 13878 13878 15075 15175 17380 17480 29417 29517 31343 31443 314483 34583 37523 37523 39614 39914 19242 22077 22177 42854 42954 45286 445386 47239 47239 49743 49843 53152 53152 5757676 24874 24974 26893 26993 29516: 31342: 31442: 10875 11921 12021 24973: contig of 3040 gap of unknown contig of 2840 gap of unknown contig of 1919 gap of unknown contig of 2424 gap of unknown contig of 1197 gap of unknown contig of 2205 gap of contig gap of contig gap of contig gap of contig gap of contig gap of contig gap of contig gap of contig contig gap of contig gap of contig gap of contig gap of contig gap of gap of contig contig contig contig contig f unknown g of 1835 f unknown
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Search completed: November Job time: 3364.8 secs
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568: gap of unknown length
458: gap of unknown length
738: contig of 3990 bp in length
5892: contig of 4280 bp in length
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5892: contig of 3144 bp in length
5892: contig of 3140 bp in length
12330: contig of 5248 bp in length
86339: contig of 5248 bp in length
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99025: contig of 4086 bp in length
99025: contig of 4686 bp in length
103329: gap of unknown length
103329: contig of 63740 bp in length
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103329: contig of 63780 bp in length
103329: contig of 6906 bp in length
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11567: contig of 6906 bp in length
125612: contig of 6906 bp in length
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131195: contig of 8483 bp in length
131195: contig of 8483 bp in length
131295: gap of unknown length
131395: gap of unknown length
131395: gap of unknown length
13149661: gap of unknown length
131662: contig of 7045 bp in length
1317964: gap of unknown length
131893: gap of unknown length
1319140: gap of unknown length
131915: contig of 8902 bp in length
1319164: gap of unknown length
1319164: gap of unknown length
1319164: gap of unknown length
131916562: contig of 8902 bp in length
131916563: contig of 8902 bp in length
1319173: gap of unknown length
131918: gap of unknown length
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131918: gap of unknown length
132918: gap of unknown length
13292: gap of unknown length
13303: gap of unknown length
13404: gap of unknown length
13504: contig of 8902 bp in length
136064: contig of 8902 bp in length
13704: gap of unknown length
13808: gap of unknown length
13909: contig of 8731 bp in length
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## SUMMARIES

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| 53.4              | 354.4              | 54.4               | 60.6               | 63.8               | 65.4               | Query<br>Score Match        |
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| AL538562          | в1913989           | BG753617           | BI917149           | BQ668420           | BI919074           | ID                          |
| AL538562 AL538562 | BI913989 603180565 | BG753617 602732803 | BI917149 6031815/1 | BQ668420 AGENCOURT | BI919074 603180881 | Description                 |

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| 9/3 8/4                                      | , co co co c                                                                           | 227.8<br>227.8<br>222.2<br>215.8<br>212.2<br>203.6                 | 505                                                      |                                                                      | 352.8<br>341.8<br>341.8<br>336.4<br>331.3<br>321.6<br>321.6<br>321.6<br>321.4<br>321.4<br>314.4<br>314.4<br>314.4<br>314.2                                                                                                                                            |
| $\omega \omega \omega \omega \omega$         | , ω ω ω υ                                                                              | 1578511                                                            | ~ ~ ~ ~ ~ ~ ~ ~                                          | 10 10 1- 01 01 01                                                    | 95.<br>92.<br>92.<br>92.<br>92.<br>6<br>92.<br>6<br>89.<br>7<br>92.<br>88.<br>7<br>9.<br>7<br>9.<br>7<br>9.<br>7                                                                                                                                                      |
| 1272<br>480                                  | 541<br>541<br>541                                                                      | 673<br>784<br>311<br>395                                           | 352<br>540<br>890<br>878<br>775                          | 788<br>664<br>1490<br>570<br>383                                     | 698<br>690<br>1010<br>704<br>705<br>850<br>340<br>654<br>948<br>976                                                                                                                                                                                                   |
| 14<br>14<br>10                               | 14                                                                                     | 10<br>12<br>13                                                     | 9<br>12<br>12                                            | 9<br>110<br>9<br>9<br>9                                              | 13<br>113<br>113<br>110<br>110<br>113<br>112<br>114<br>114<br>114                                                                                                                                                                                                     |
| BM981455<br>BM981442<br>BQ224442<br>BE336607 | H1/169<br>A1767459<br>BM974798<br>BW717094                                             | BB644064<br>BG704350<br>AU050830<br>AW417548<br>BI544761<br>R20352 | AA349894<br>AW954549<br>BF125425<br>BF125664<br>AU035640 | AU080744<br>BB652617<br>AK012406<br>B1831919<br>AI614768<br>AU067460 | B1554034<br>B1669845<br>B345141<br>BE386060<br>B1823714<br>BG753353<br>BE736301<br>H06756<br>BE746601<br>B0942438<br>B0942438<br>B0942438<br>B0942438<br>B0942438                                                                                                     |
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| found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLAM11618 row: a column: 21 High quality sequence start: 4 High quality sequence stop: 702. Location/Qualifiers                                                                                               |
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| Email: cgapbs-r@mail.nin.gov Tissue Procurement: Life Technologies, Inc. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be |
| JOURNAL Unpublished (1999) MMENT Contact: Robert Strausberg, Ph.D.                                                                                                                                                                                                                          |
| AUTHORS NIH-MGC http://mgc.nci.nin.gov/. TITLE National Institutes of Health, Mammalian Gene Collection (MGC)                                                                                                                                                                               |
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| ŪRCE human.<br>ORGANISM Homo sapiens                                                                                                                                                                                                                                                        |
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| mRNA sequence.<br>BI919074                                                                                                                                                                                                                                                                  |
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                                        Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                     NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
                                                                             Contact: Robert Strausberg, Ph.D
                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                             AGENCOURT_8301981 NIH_MGC_102 Homo sapiens cDNA clone IMAGE:6275121
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cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism-"Homo sapiens"
/db_xref-"taxon:9606"
/clone-"IMAGE:5244956"
/clone_lib-"NIH_MGC_121"
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 747)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                    EST.
                                                                                                                                                                                                                                     603181571F1 NIH_MGC_121 Homo sapiens cDNA clone IMAGE:5245648 5',
                                                                                                                     Homo sapiens
                                                                                                                                                                                      BI917149.1 GI:16181111
                                                                                                                                           human.
                                                                                                                                                                                                                                 mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="epidermoid carcinoma, cell line"
/lab_host="DH10B (phage-resistant)"
/note="Organ: sallvary gland; Vector: pOTB7; Site_1: xhoI;
Site_2: ECORI; cDNA made by Oligo-dT priming.
Directionally cloned into EcORI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-CDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library."

96 a 314 c 285 g 191 t l others
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/db_xref="taxon:9606"
/clone="IMAGE:6275121"
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Pred. No. 3.2e-102;
0; Mismatches 4;
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mRNA sequence.
BG753617
BG753617.1 GI:
EST.
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Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
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Contact: Robert St
                                                                        BG753617 670 bp r 602732803F1 NIH_MGC_43 Homo sapiens
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/db_xref="taxon:9606"
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/clone_lib="NIH_MGC_121"
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98.4%;
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IMAGE:4876392 5',
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                      CCACCATCCT 369
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Tissue Procurement: ATCC
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National Institutes of Health, Mammalian Gene
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/tissue_type="normal pigmented retinal epithelium"
/lab_bost="PH10B (phage_resistant)"
/note="Organ: eye: Vector: pOTB7; Site_1: XhoI; Site_2: EcoRI; cDNA made by Oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5; adaptor: GGCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH_MGC Library. | "
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Pred. No. 2.1e-99;
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                 CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCANGCCAT-GGCTCCGCA
                                                                                                                                                                                                                                                                                                                     GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCAC
CGGGCCAAGGCCCAAGAAAAGGGGAAGTTCTGCCTCGGCCCTCAGGCCATGGGCTCCGGA
                                                                                         CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAGCACCCCCTCTTTGTAA
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                                                                        CTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCCTCTTTGTAA
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Tissue Procurement: Life Technologies, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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National Institutes of Health, M
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603180565F1 NIH_MGC_121
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://imagge.llnl.gov
plate: LLAM11617 row: a column: 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note-"organ: brain; Vector: pCMV-SPORT6; Site_1: Not1; Site_2: ECORV (destroyed); RNA source anonymous pool of 3 fetal brains, female age 20 weeks, female age 24 weeks, and male age 26 weeks. Library is oligo-dT primed and directionally cloned (ECORV site is destroyed upon cloning). Average insert size 1.7 kb, insert size range 0.7-3.5 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 017. Note: this is a NIH_MGC Library."
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/clone_lib="NIH_MGC_121"
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BP 191 91006 EVRY cedex - France
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="Organ: Fetal brain; Vector: pCMVSPORT 6; 1st strand /note="Organ: Fetal brain; Vector: pCMVSPORT 6; 1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and Eco RV sites of the pCMVSPORT 6 vector. Library was constructed by Life Technologies. Contact: Feng Liang Life Technologies, a division of Invitrogen 9800 Medical Center Drive Rockville Maryland 20850, USA Fax: (1) 301 610 8371 Email:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 http://fulllength.invitrogen.com"
285 c 257 g 199 t
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/clone_lib="LTI_FL013_FBrn1"
/clone_lib="LTI_FL013_FBrn1"
/dev_stage="pooled tissue from post conception fetuses (20 week, 24 week and 26 week)"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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TCTGCTCCCCAGGGAAACTGAACTCAGTTTGCATCAGCTGCTGCAACACCCCTCTTTGTA 299
                                                                                                 ACCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCT
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Tissue Procurement: David N. Louis, M.D.

CDNA Library Preparation: Life Technologies, Inc.

CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can be
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Plate: LLAM11784 row: d column: 08
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National Institutes of Health, Mammalian
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/note="Organ: brain; Vector: pCMV-SPORT6; Site_1: NotI;
Site_2: SalI; Cloned unidirectionally. Primer: Oligo dT
Average insert size 2.3 kb. Constructed by Life
Technologies. Note: this is a NCI_CGAP Library."
a 231 c 204 g 132 t
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/clone="IMAGE:5309527"
/clone_11b="NCI_CGAP_Brn67"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /tissue_type="anaplastic oligodendroglioma with 1p/19q
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98.6%;
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Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                    Score 352.8; DB 1 pred. No. 6.9e-99;
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                                                                             GGTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCA
                                                                                               -GTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCA 119
CGGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGT
                                   CGGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGT 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tissue Procurement: Miklos Palkovits, M.D., Ph.D. cDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki Toshiyuki and Piero Carninci (RIKEN) cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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BI669845
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (1999)
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National Institutes of Health, Mammalian Gene Collection (MGC)
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Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (bases 1 to 690)
                                                                                                                                                                                                                                                                                                                                       127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                quality sequence stop:
                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                  /note="Organ: brain; Vector: pBluescriptR (modified pBluescript KS+); Site_1: BamHI; Site_2: SalI-xhoI (gtcga); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library. a 231 c 202 g 130 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="hypothalamus"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone="IMAGE:5312889"
/clone_lib="NIH_MGC_96"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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                                                                                                                                                                                                                                                       Score 343.4;
Pred. No. 5.9
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287 AGTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCA 346
                                                                                   227
                                         60
                                                                                                                                                                                         Local
                                                                                                          GGCAACTTTTTGCGGATTGTTCTTGCTTNC-AGGCTTTGCGCTGCAAATCCAGTGCTACC 59
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                                                                                   GGCAACTTTTTGCGGATTGTTCTTGCTTCCAAGGCTTTGCGCTGCAAATCCAGTGCTACC 286
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                      AGTGTGAAGAATTCCAGCTGAACAACGACTGCTCCTCCCCCGAGTTCATTGTGAATTGCA 119
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N. Louis, M.D.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 1010)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1999)
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Plate: LLAM9422 row: d column: 05
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                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                            /lab_host="DH10B (TI phage-resistant)"
/note="Organ: brain; vector: pCMV-SPORT6; Site_1: Not1;
Site_2: Sall; Cloned unidirectionally, Primer: Oligo dT.
Average insert size 2.3 kb. Constructed by Life
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                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4153684"
/clone_lib="NCI_CGAP_Brn67"
                                                                                                                                                                                                                                                                                                                                                                                                                        /tissue_type="anaplastic oligodendroglioma with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ocation/Qualifiers
                                                                                                                                                                                       92.6%;
                                                                                                                                                                     0;
                                                                                                                                                               Score 341.8; DB LZ;
Pred. No. 2.4e-95;
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                                                                                                                                                                                                           Length 1010;
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                 31 AGGCTTTGCGCTGCAAATCCAGTGCTACCAGTGTGAAGAATTCCAGCTGAACAACGACTG 90
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                                                                                              337;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTP/DTP
cDNA Library Preparation: Ling Hong/Rubin Laboratory
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov
Plate: LLCMA85 row: 1 column: 14
High quality sequence stop: 593.
Location/Qualifiers
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1 (bases 1 to 593)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Unpublished (1999)
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BE386060
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                                                                                                                      Similarity
                                                                                              Conservative
                                                                                                                                                                                                             /tissue_type="melanotic melanoma"
/lab_host="DH10B (phage-resistant)"
/note="Organ: skin: Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Organ: skin: Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Organ: skin: Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Organ: skin: Vector: poTB7; Site_1: XhoI; Site_2: Directionally
/cloned into EcoRT/XhoI sites using the following 5;
/cloned into EcoRT/XhoI sites using the following 5;
/cloned into EcoRT/XhoI sites using the following 5;
/cloned into EcoRT/XhoI sites using the following 5;
/cloned into EcoRT/XhoI sites using the following in the laboratory of Gerald M. Rubin (University of the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit
/cStratagene) and Superscript II RT (Life Technologies)."

39 a 166 c 183 g 105 t
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/clone_lib="NIH_MGC_20"
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                                                                                                                      91.2%;
99.4%;
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                                                                                                                      Score 336.4; DB 1 Pred. No. 8.2e-94;
                                                                                                 Mismatches
                                                                                                                                             DB 10;
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SOURCE
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AUTHORS
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            Query Match
Best Local Similarity 97.5
Matches 356; Conservative
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                                                                                                                                                                                                                                                                                                                                                                      found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov Plate: LLAMI1453 row: a column: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
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1 (bases 1 to 704)
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BI823714
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National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                            quality sequence stop:
                                                                                     /note="Organ: pooled brain, lung, testis; vector: pCMV-SPORT6; Site_1: Not1; Site_2: EcoRV (destroyed); RNA source anonymous pool of 6 male brains, age range 23-27; 1 male lung, age 27; and 1 male testis, age 69. Library is oligo-dT primed and directionally cloned (EcoRV site is destroyed upon cloning). Average insert size 1.8 kb, insert size range 1-3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 021. Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5181585"
/clone_lib="NIH_MGC_115"
                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                     /lab_host="DH10B"
                                89.78;
                 Score 331; DB 13;
Pred. No. 4.4e-92;
0; Mismatches 7
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BASE COUNT
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1 (bases 1 to 705)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               found through the I.M.A.G.E. Consortium/LLNL http://image.llnl.gov plate: LLCM1756 row: o column: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (1999)
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602731780F1 NIH_MGC_43 Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         quality sequence stop: 481
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/clone_lib="NH_MCC_43"
/tlissue_type="normal pigmented retinal epithelium"
/lab_host="PH10B (phage-resistant)"
/note="Organ: eye: Vector: pOTB7; Site_1: xhoI; Site_2: ECORI; cDNA made by oligo-dT priming. Directionally cloned into ECORI/XhoI sites using the following 5; adaptor: GCCACGAG(G). Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MCC Library. | "
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/clone="IMAGE:4875499"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mRNA sequence.
BE736301
                                                                                                                                                                                                   CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                            Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                     Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                     NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, M
                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                      BE736301.1 GI:10150293
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BE736301 850 bp mRNA linear EST 15-SEP-2000 601307087F1 NIH_MGC_39 Homo sapiens cDNA clone IMAGE:3641660 5',
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                                                                                                                                                                           http://image.llnl.gov
Plate: LLCM347 row: m column:
                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                          High quality sequence stop: 662.
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/tissue_type="adenocarcinoma"
/lab_host="0H10B (phage-resistant)"
/note="Organ: pancreas; Vector: pOTB7; Site_1: XhoI;
/note="Organ: pancreas; Vector: pOTB7; Site_1: XhoI;
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Directionally cloned into EcoRI/XhoI sites using the
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                                                                                   /clone="IMAGE:3641660"
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4

Source: IMAGE Consortium, LLNL;
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further i
Insert Length: 1737 Std Error: 0.00
Seq primer: M13RP1

further information

contact the

Email: est@watson.wustl.edu Insert Size: 1737 High quality sequence stops: 312

Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. I Tel: 314 286 1800 Fax: 314 286 1810

Louis, MO

Unpublished (1995) Contact: Wilson RK

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Best Local Similarity
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yl83b07.rl Soares infant brain 1
IMAGE:44619 5', mRNA sequence.
H00756
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

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                                                                                                                                                                                                                                                                                                                       -CCAAGAAAAGGGGAAGTTCTGCCTTGGCCCTCAGGC 336
                                                   NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 654)
                                                                                                                                                       Homo sapiens
                                                                                                                                                                                 BE746601
BE746601.1 GI:10160593
EST.
                                                                                                                                                                                                                       mRNA sequence.
                                                                                                                                                                                                                                    601580186F1 NIH_MGC_9 Homo sapiens
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cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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Location/Qualifiers
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/dev_stage="73 days post natal"
/lab_host="DH10B (ampicillin re
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/clone="IMAGE:44619"
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/db_xref="GDB:417160"
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98.8%;
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Pred. No. 2.7e-89;
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                                                                                                                                                                                                                                                                                                                                              CGGTGAACGTTCAAGACATGTGTCAGAAAGAAGTGATGGAGCAAAGTGCCGGGATCATGT
                                                                                 ACGGGCCAAGGCCCAAGACAAGGGGAAGTTCTGCCTCGGCC
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Location/Qualifiers
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/clone_lib="NIME_MGC_9"
/clone_lib="NIME_MGC_9"
/tissue_type="denocarcinoma cell line"
/tissue_type="denocarcinoma cell line"
/lab_host="DHIOB (phage-resistant)"
/lab_host="DHIOB (phage-resistant)"
/note="Organ: ovary: Vector: pOTB7; Site_1: XhoI; Site_2:
/note="Organ: ovary: Vector: pOTB7; Site_1: XhoI; Site_2:
/cloned into EcoRI/XhoI sites using the following 5;
cloned into EcoRI/XhoI sites using the following 5;
adaptor: GGCACGAG(G). Size-selected >500bp for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
21 a 219 c 187 g 126 t 1 others
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/db_xref="taxon:9606"
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Maximum DB seq length: 2000000000
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|                    | 577      | 23 | Ω    |      | 21  | 44       |
| Human prostate exp | ABV05091 | 23 | œ    |      | 21  | ü        |
| Human breast cance | AAL09957 | 22 | æ    |      | 21  | 2        |
| Human cervical can | AAH69474 | 22 | ഗ    |      | 21  | ï        |
| Human polynucleoti | 081      | 24 | 2049 |      | 22  | 0        |
| Human colon cancer | AAH34281 | 22 | N    |      | 22  | 39       |
| Arabidopsis cDNA e | 537      | 24 | æ    |      | 22  | 8        |
| Human polynucleoti | 9048     | 24 | 1329 |      | 22  | 37       |
| nucle              | 25       | 22 | 4    |      | 22  | 36       |
| Human prostate exp | 833      | 23 | 377  | 3.5  | 22  | 5        |
| Human head and nec | ABK53839 | 24 | ū    |      | 22  | 4        |
| tate               | S        | 23 | 1539 |      | 23  | ü        |
| Human prostate exp | ABV23163 | 23 | 1539 |      | 23  | ະ        |
| Human breast cance | AAL10167 | 22 | 377  |      | 23  | 3        |
| Human nervous syst | æ        | 22 | 263  |      | 23  | ö        |
| Human SNP oligonuc | ~        | 22 | 51   |      | 25  | 29       |
| 011                | 398      | 22 | 50   |      | 26  | 8        |
| Human SNP oligonuc | 8        | 22 | 50   | 4.2  | 26  | 27       |
| Human ovarian canc | 8484     | 24 | 381  | 4.   | 92  | 6        |
| ovarian            | 793      | 24 | 558  | 9    | 120 | 8        |
|                    | 855      | 24 | 390  | 4.   | 154 | 24       |
| Human ovarian canc | 591      | 24 | 201  | 5    | 166 | 23       |
| ian                | 126      | 24 | 409  | Ν.   | 263 | 22       |
| Human ovarian canc | 943      | 24 | 349  | 49.1 | 307 | 2        |
| 5                  | 53       | 24 | 373  | 0    | 314 | õ        |
| tumour asso        | 0        | 22 | 444  | ა    | 352 | 6        |
|                    | 698      | 22 | 52   | 7.   | 359 | 8        |
| rcin               | ဒ        | 24 | 9    | 7.   | 359 | .7       |
| ovarian ca         | 0327     | 24 | 36   | 7.   | 359 | 9        |
| -                  | AAF94186 | 22 | æ    | в.   | 365 | <u>5</u> |
| cDNA encoding huma | ABK33543 | 24 | 52   | 2    | 393 | 4        |
| Ovarian carcinoma  | 34       | 24 | 61   | .7   | 486 | ω        |
| Ovarian carcinoma  | 2        | 24 | 61   | 7.   | 486 | ົວ       |
| ovarian car        | 032      | 24 | 1619 | 77.8 | 486 | _        |
| Human ovarian carc | ABT03277 | 24 | 61   | .7   | 486 | 0        |
|                    |          |    |      |      |     |          |

## ALIGNMENTS

RESULT 1
ABT03280
ID ABTC

ABT03280 standard; cDNA; 625 BP.

05-SEP-2002 (first entry)

ABT03280;

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14-NOV-2000; 2000US-0713550
03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
                                                                                                                                                                    Human; ovarian cancer; ovarian carcinoma; gene therapy; immunotherapy; cytostatic; gene; ss.
                WPI;
                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                             Human ovarian carcinoma associated coding sequence SEQ ID NO: 210.
                                 Xu J,
                                                                                                   13-NOV-2001; 2001WO-US45395
                                                                                                                    23-MAY-2002
                                                                                                                                    WO200239885-A2
                                                 (CORI-) CORIXA CORP.
                2002-500186/53.
                               Stolk JA, Algate
                                 PA,
                                Fling SP,
                                 Molesh DA
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625 625 625 1897 1897 1608 1953 1953

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ABL87898 ABT03284 ABL40352 ABQ54231 AAF22400 AAF64188 AAF93845

Human ovarian canc Human ovarian carc Ovarian carcinoma Human ovarian anti

Human ovarian carc Ovarian carcinoma

Human secreted pro Human secreted pro Human cDNA encodin

Novel ovarian cancer polypeptide and polynucleotide, useful for

Score

Length DB

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Description

ABT03280 ABL40348

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RESULT 2
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                            Ovarian carcinoma 01034C EST clone nucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides human ovarian cancer and coding sequences. The sequences can be used in treatment of ovarian cancers. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian cancer
  Ovarian cancer;
                                                         28-JUN-2002
                                                                                                               ABL40348
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                                                                                                                                                                                TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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                                                                                                               standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             625 BP; 182 A; 144 C; 182 G; 116 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 194; 197pp; English
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                                                         (first entry)
  cancer therapy; vaccine;
                                                                                                               CDNA;
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gene therapy; tumour;
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                                                                                                                                                                                                                                                                                                                                    that may be utilised in cancer therapy, for example in a vaccine or compensation or therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or companies of a cancer in a patient, for stimulating the compensation of a cancer in a patient. They are also useful for compensation of a cancer in a patient. They are also useful for their compensation of determining the presence of a cancer in a patient.

CC a patient and for determining the presence of a cancer in a patient.

CC The isolated polynucleotides of the invention are useful for their complimatory in the complementary stretches of the entire desired gene or gene fragments, and for designing and complementary in the complementary stretches complementary in the complementary stretches in tumour cells. Polypeptides and polynucleotides of the convention are also useful in recombinant DNA molecules to direct compression of a polypeptide in appropriate host cells. The current consequence represents the ovarian carcinoma old34C EST clone nucleotide consequence.
                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 625; Conserv
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15-AUG-2000;
07-SEP-2000;
14-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (XUJJ/)
(STOL/)
(ALGA/)
                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                              sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim la; Page 125;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ovarian tumour polypeptide and polynucleotide useful in diagnosis, prevention and/or treatment of cancer, especially ovarian cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI;
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  GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
               GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTTGGCCGCAGC
                                                                     CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAAGGGGAG
                                                                                                             CAGCGACAGGCGGCAGCACAGCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACAC
                                                                                                                                                                                GCCCTCTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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                                                                                                                                                                                                                                     AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
                                                       CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
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) STOLK J A.
) ALGATE P A.
) FLING S P.
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The present invention describes a composition (I) comprising: carriers and immunostimulants; and a polypeptide (II) of a ovarian tumour polypeptide encoded by a polynucleotide (III) having a cDNA sequence (S1) from the 1091 nucleotide sequences as given in ABL77023 to ABL87934, (III) encoding (II) having a sequence (S2), a T cell population of (II), or antigen presenting cells that express (II). (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to (S1) can be used for detecting ovarian cancer in a patient's biological sample preferably serum or ovarian tissue. The method comprises contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to patient wather the amount to a predetermined cutoff value and thereby detecting ovarian cancer in the
                                                                                                                                                                                                                                                                                                                                                                                                                     Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing polypeptide
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                                                                                                                                                                                                                                                                                                                                                                    Claim 1;
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(first entry)

ABT03284 standard;

CDNA; 1897

Human ovarian carcinoma associated coding

cytostatic;

gene;

Human; ovarian cancer;

ovarian carcinoma; gene therapy; immunotherapy;

sequence SEQ

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Pred. No. 5.2e-239;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-NOV-2000; 2000US-0713550; 03-APR-2001; 2001US-0825294; 02-OCT-2001; 2001US-0970966;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1897 BP; 435 A; 521 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; Page 196; 197pp;
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                                                    TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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RESULT 5
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                                                                                                 that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their
                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
14-NOV-2000;
                 ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polypuclectides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current
                                                                                                                                                                                                                                                                                                                                                                                                 (XUJJ/)
(STOL/)
(ALGA/)
           sequence
                                                                                                                                                                                                                   The invention relates to ovarian tumour polynucleotides and polypeptides
                                                                                                                                                                                                                                              Claim la; Page 127-128; 131pp; English
                                                                                                                                                                                                                                                                       Ovarian tumour polypeptide and polynucleotide useful in diagnosis, prevention and/or treatment of cancer, especially ovarian cancer
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) STOLK J A.
) ALGATE P A.
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           represents the ovarian carcinoma 01034C/0591S
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2000US-0640173.
2000US-0656668.
2000US-0713550.
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/product= "Ovarian carcinoma protein Ol034C/O591S"
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                                Human; ovarian antigen; ovary; ovarian; breast; cancer; tumour; ovarian cancer; breast cancer; tumour; reproductive system disorder; infertility; pregnancy disorder; anovulation; polycystic ovary syndro; PCOS; ovarian cyst; dysmenorrhoea; endocrine disorder; infection; inflammatory condition; inmune disorder; blood disorder; cardiovascular disorder; respiratory disorder; neurological disorder; gastrointestinal disorder; urinary system disorder; drug screening; gene therapy; chromosome mapping; forensic analysis;
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                     gene therapy; chromosome mapping;
antibody preparation; cytostatic;
                                                                                                                                                      Human ovarian antigen HVCAG04 cDNA,
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          gynaecological;
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          reproductive;
                        immunomodulatory; neuroprotective
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                                                                                                                                                       SEQ ID NO:111.
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AGTTCTCCTTGCAGAGGACTGGCGCGCGGACGCGAGAGAGCAACGGGCGCTGCACAAAGCG

AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG

925

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Query Match Matches

Local

Similarity

99.8%;

Score 555; DB 24 Pred. No. 3e-211; Mismatches

DB 24;

Length Indels

1608; 0;

Gaps

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Conservative

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1045

CAGCGACAGGCGGCACAGCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACAC

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120 1044

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CC ABP4328) and to cDNAs encoding them (ABC54111-ABC56305), and also CC encompasses polypeptides 90% identical and polynucleotides 95% identical to the sequences of the invention. The invention additionally relates to recombinant vectors and host cells comprising human ovarian antigen CC polynucleotides, antibodies against human ovarian antigen and the use covarian antigen polynucleotides and polypeptides in diagnosing, cc treating, prognosing or preventing various ovary and/or breast cancer, and disorders. Such conditions include ovarian cancer and breast cancer, and disorders. Such conditions include ovarian cancer and breast cancer, and cdisorders (e.g., infertility, disorders of pregnancy, anovulation, polycystic ovary syndrome, ovarian cysts, and dysmenorrhoea), endocrine cdisorders, infections (e.g., chlamydia, HIV, toxoplasmosis, and toxic shock syndrome), inflammatory conditions (e.g., mastitls, oophoritis and cysinitis), immune disorders (e.g., congenital and acquired immunodeficiencies, autoimmune oophoritis, systemic lupus erythematosus), blood-related disorders (e.g., anaemia), cardiovascular disorders, crespiratory disorders, neurological disorders, gastrointestinal disorders of polynucleotides may also be used in screening for compounds which conjugate therapy, chromosome mapping, in the colypeptides may also be used in screening for compounds which colypeptides may be used as food additives or to prepare antibodies useful in disease diagnosis, drug targeting and phenotyping. The present conjugate colons in the present colons of the invention.
                                      specification, but was obtained in electroni
at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Isolated nucleic acid molecules encoding novel ovarian polypeptides, useful in the prevention, treatment and diagnosis of cancer (e.g. ovarian cancer), immune disorders, cardiovascular disorders and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-2000;
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Sequence 1608 BP; 402 A; 402 C; 417 G; 381 T; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1;
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                                                                                     sequence data
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               relates to 2175 novel human ovarian antigens (ABP41054-to cDNAs encoding them (ABQ54131-ABQ56305), and also
                                                               ence data for this patent did not form part of the printed but was obtained in electronic format directly from WIPO
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other;
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Isolated nucleic acid molecule encoding a human secreted protein used in preventing, treating or ameliorating a medical condition  $% \left( 1\right) =\left( 1\right) +\left( 1\right) +$ 

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            P-PSDB;
                      WPI; 2000-647420/62
                                                                                            09-APR-1999; 99US-0128694.
20-JAN-2000; 2000US-0176931.
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                                         Ruben SM,
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                                                             GENOME SCI INC
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ytostatic; cardiant; vasotropic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       present invention.
GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                            GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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or diagnose various diseases and disorders including, autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. Alzheimer's disease, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin ageing due to sumburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to
                                                                                                                                                                                                                                                                                                                                                                       The proteins and polynucleotides, their agonists and antagonists have activities dependent on the tissues and cells in which they are expressed, examples of these activities include, immunosuppressive; antiarthritic; antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; opthalmalogical; and vulnerary. The proteins, polynucleotides, agonists and antagonists can be used to treat or detect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted proteins AAB75506 - AAB75554 are encoded by polynucleotic sequences AAF64176 - AAF64224. The specification includes amino acid sequences AAB75555 - AAB75606 which represent fragments of the human secreted proteins, and protein sequences with which they share homology secreted proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid molecules encoding human secreted proteins, used preventing, treating or ameliorating a disorder, e.g. Alzheime Parkinson's diseases and cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUN-2000; 2000WO-US14973
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; secreted protein; immunosuppressive; antiarthritic; antirheumatic;
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                                          chemotaxis. The
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Matches 605
                                                                      Human cDNA encoding a membrane or secretory protein clone PSEC0181
                                                                                                                                 AAF93845 standard;
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                          Homo sapiens
                                            Human; secretory protein; rheumatoid arthritis; dia
                                                                                            23-MAY-2001
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11-JAN-2000;
02-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein/membrane protein expression. The nucleic acids and complementary sequences may also be used as DNA probes in diagnostic assays (e.g. polymerase chain reactions (PCR)) to detect and quantitate the presence of similar nucleic acid sequences in samples. They may also be used to study the expression and function of secretory proteins/membrane polypeptides and their role in metabolism. The polypeptides may be used as antigens in the production of antibodies against them and in assays to identify modulators (agonists and antagonists) of expression and activity. The antibodies and antagonists may also be used as therapeutic agents to down regulate expression and activity. The antibodies agents for detecting the presence of the polypeptides in samples (e.g. by enzyme linked immunosorbant assay (ELISA). Examples of diseases which may be treated include rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AABB8317 - AAB88419. Included in the invention are primers AAF93917 - AAF94295 and AAF62232 - AAF62233 which are used to isolate to DNA sequences of the invention. The invention also includes methods in the production of antibodies directed against the proteins, and cDNA sequences, which can be used in vaccines. The polynucleotide sequences can be used in gene therapy. The polynucleotide sequences can be used in gene therapy. The polynucleotide sequences and the proteins they encode may be used in the prevention, treatment and diagnosis of diseases associated with inappropriate secretory
                                                          1530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to nucleic acid sequences AAF93744 - AAF93916 which encode human secretory or membrane proteins represented by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acids encoding secretory proteins/membrane progene therapy or as candidate target molecules in drug
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                                                                                                                                                                                                                            CAGGCGGCAGCACACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCCTGTA 186
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                   CCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCA
                                                                                                                                                                     CAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGAT
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ruq development –
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                                                                                                                                                                        Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
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                                                 Sequence 1619 BP;
                                                                                     The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                      Claim 2;
                                                                                                                                                                                                                             WPI; 2002-500186/53
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                                                                                                                                                                                                                                                    Stolk JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ovarian cancer; ovarian carcinoma; gene therapy;
                                                                                                                                      Page 189-190; 197pp; English.
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Score 486; DB 24;
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03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
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cancer polypeptide and polynucleotide, useful for presence of ovarian cancer in a patient, and in
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(first entry)

cDNA; 1619

cancer therapy; vaccine; gene therapy; tumour;

cancer;

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Query Match
Best Local Simi
Matches 606;
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99.8%;
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Pred. No. 7.8e-184;
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Homo

sapiens

US2002004491-A1

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The invention relates to ovarian tumour polynucleotides and polypeptides CC that may be utilised in cancer therapy, for example in a vaccine or CC gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or CC expanding T-cells specific for a tumour protein, and for inhibiting the CC development of a cancer in a patient. They are also useful for CC stimulating an immune response in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient. CC The isolated polynucleotides of the invention are useful for their CC ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and CC preparing ribozyme molecules for inhibiting expression of tumour CC invention are also useful in recombinant DNA molecules to direct CC expression of a polypeptide in appropriate host cells. The current CC sequence represents the extended CDNA sequence of ovarian carcinoma CC isolate 57887 given in record ABL48956.
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) STOLK J A.
) ALGATE P A.
) FLING S P.
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t of
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gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient, and for treating a cancer in a patient.
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                  Ovarian tumour polypeptide and prevention and/or treatment of
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STOLK J A
ALGATE P.
FLING S P
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131pp;
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                   polynucleotide useful in diagnosis, cancer, especially ovarian cancer
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   breast cancer; prostate tumour; rectal tumour; liver tumour;
                                             cDNA encoding human PRO protein, Seq ID No
                                                                                                                                    ABK33543 standard; cDNA; 1524
                 Human; secreted protein; PRO; tumour; lung cancer;
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99.8%;
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Matches

Similarity

100.0%; 62.9%;

Score 393; DB 24; Pred. No. 7e-147; Mismatches

Length 1524;

Conservative

0,

0

Indels

0;

0

60

Sequence

1524 BP; 321 A; 433 C; 435 G; 335 T; 0 other;

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26-JUL-2000;
28-JUL-2000;
23-AUG-2000;
24-AUG-2000;
11-SEP-2000;
10-NOV-2000;
28-NOV-2000;
20-DEC-2000;
20-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-JUL-2000;
25-JUL-2000;
25-JUL-2000;
                        The PRO polypeptides are useful for diagnosing tumours, especially lung cancer, colon cancer, breast tumour, prostate tumour, rectal tumour or liver tumour. The PRO polypeptides are useful for stimulating the proliferation of, or gene expression, in pericyte cells, for stimulating the proliferation or differentiation of chondrocyte cells, for stimulating the release of tumour necrosis factor-alpha from human blood, for stimulating or inhibiting the proliferation of normal human dermal fibroblast cells. The PRO polypeptide may also be used as molecular weight markers and for tissue typing. The PRO nucleic acids have applications in molecular biology, including use as hybridisation probes, and in chromosome and gene mapping. ABK33536-ABK33657 represent human PRO protein coding sequences of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     pericyte cell proliferation; chondrocyte cell proliferation; tumour necrosis factor-alpha; gene; ss.
                                                                                                                                                                                                                  The invention relates to one hundred and twenty two nucleic acids encoding PRO polypoptides. The sequences of the 122 PRO polypoptides, encode human secreted proteins. The PRO nucleic acids, polypoptides, agonists and antagonists are useful for treating a PRO related disorder.
                                                                                                                                                                                                                                                                                                                             One hundred and twenty two nucleic acids encoding PRO polypeptides, useful for treating a PRO related disorder and for diagnosing tumours such as lung cancer, colon cancer, breast tumour, prostate tumour, retumour or liver tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Grimaldi JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-JAN-2002
                                                                                                                                                                                                                                                                                               Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Baker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                           2002-172001/22.
DB; AAU83599.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KP,
                                                                                                                                                                                                                                                                                                <u>ب</u>
                                                                                                                                                                                                                                                                                                Figure 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Desnoyers L,
C, Gurney AL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000WO-US23522
2000WS-00000D
2000WS-00000DP
2000WS-US30873
2000WS-US3646P
2000WS-US32678
2000WS-US32678
2000WS-US34956
2001WS-US34956
2001WS-US34956
2001WS-US34956
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                                                                                                                                                                                                                                                                                                359pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gerritsen ME, Goddard A, Godowski Smith V, Stephan JF, Watanabe CK,
                                                                                                                                                                                                                                                                                                                                              prostate tumour, rectal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Godowski PJ;
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RESULT 15
AAF94186/c
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which encode human secretory or membrane proteins represented by AAB88317 - AAB88419. Included in the invention are primers AAF93917 - AAF94295 and AAF62232 - AAF62235 which are used to isolate the CDNA sequences of the invention. The invention also includes methods for the production of antibodies directed against the proteins, and cDNA sequences, which can be used in vaccines. The polynucleotide sequences can be used in gene therapy. The polynucleotide sequences and the proteins they encode may be used in the prevention, treatment and diagnosis of diseases associated with inappropriate secretory
                                                                                                                                                                                                                                                                                                                                                            08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
                                                                                                                                                This invention relates to nucleic acid sequences AAF93744 - AAF93916
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAF94186 standard; DNA; 587
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                                                                                                                                                                                 Claim 5; SEQ ID 620; 609pp + CD ROM; English.
                                                                                                                                                                                                                             Nucleic acids encoding secretory proteins/membrane proteins, useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; secretory protein; membrane protein; vac rheumatoid arthritis; diabetes; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-MAY-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1192
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                                                                                                                                                                                                                 therapy or as candidate target molecules in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGGGCAGACACGAGCAGGGGGGGGGAGACAGAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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                                                                                                                                                                                                                                                                                              Isogai T,
                                                                                                                                                                                                                                                                                                                                                            2000JP-0118775
2000JP-0183766
                                                                                                                                                                                                                                                                                                                                                                                              99JP-0194179.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for DNA encoding secretory/membrane protein SEQ ID
                                                                                                                                                                                                                                                                                                  Nishikawa
                                                                                                                                                                                                                                                                                                H,
                                                                                                                                                                                                                                                                                              Kawai Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                393
                                                                                                                                                                                                                                                                                              Sugiyama
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      vaccine; gene therapy,
                                                                                                                                                                                                               drug
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                                                                                                                                                                                                                 development
                                                                                                                                                                                                                                                                                              Hayashi
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                                                                                                                                                                              cytostatic;
                                                                                                               WO200239885-A2
                                                                                                                                               Homo sapiens.
                                                                                                                                                                                           Human; ovarian
                                                                                                                                                                                                                              Human ovarian carcinoma associated coding sequence SEQ ID NO:
                                                                                                                                                                                                                                                                05-SEP-2002
                                                                                                                                                                                                                                                                                                                               ABT03279 standard; cDNA; 1362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 587 BP; 117 A; 162 C; 140 G; 162 T; 6 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein/membrane protein expression. The nucleic acids and complementary sequences may also be used as DNA probes in diagnostic assays
                                                                                                                                                                                                                                                                                                                                                                                                                                           551
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           117
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                                                                                                                                                                                                                                                                                                                                                                                                              CAAACCCGGGGGAGTTGGCTGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG
                                                                                                                                                                                                                                                                                                                                                                                                                             CAAACCCGGGGGAGTTGGCTGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATT 490
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGGTTTGGCCGCAGCCATGATCCTC 298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTC 310
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATT 118
                                                                                                                                                                              gene;
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                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                           cancer;
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                                                                                                                                                                                               ovarian carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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No. 1e-135;
                                                                                                                                                                                                                                                                                                                               ВP
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                                                                                                                                                                                             gene therapy; immunotherapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        <u>..</u>
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14-NOV-2000; 03-APR-2001;

13-NOV-2001; 2001WO-US45395

23-MAY-2002

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RESULT 17
ABL40347/c
ID ABL40347 standard;
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                                                                                                                                                                                                                                                                                               1155
                                                                                                                                                                                                                                                                                                                           1215
      US2002004491-A1
                                                         Homo
                                                                              Ovarian
                                                                                            Ovarian
                                                                                                           28-JUN-2002
                                                                                                                         ABL40347;
                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1362 BP;
                                                                                                                                                                                                                                                                                                                                                                                                    ō,
                                                                                                                                                                                                                                                                                                                                                                                                       The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-OCT-2001; 2001US-0970966
                                                                                                                                                                                          301
                                                                                                                                                                                                                       241
                                                                                                                                                                                                                                                    181
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                                                                                                                                                                                                                                                                                                                                                                       Match
                                                                                                                                                                                                                                                                                                                                                                                                   the invention.
                                                                                                                                                                         sapiens
                                                                                                                                                                                                     GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
                                                                                                                                                                                                                                  CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAACGTCTCCGAGAAGGGGAG
                                                                                                                                                                                                                                                                CAGCGACAGGCGGCACCACCACCTGCACGACACCCGGCGGAAACTGCTGCGAGGACAC
                                                                                                                                                                                                                                                                                                     GCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                                                                                                                                                                                            GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                       2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Stolk JA, Algate
                                                                              cancer;
                                                                                            carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 193; 197pp; English.
                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                          (first entry)
                 /*tag= a
/product=
                                                                          cancer therapy; vaccine; gene therapy; tumour; cancer;
                                 Location/Qualifiers
1..1362
                                                                                           GPR39 cDNA
                                                                                                                                                                                                                                                                                                                                                                                    263 A; 435
                                                                                                                                                                                                                                                                                                                                                             57.4%;
100.0%;
                                                                                                                                       1362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PΑ,
                 "Ovarian carcinoma
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                  protein GPR39
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10-SEP-1999;
01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
14-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the overten carriers are page.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene therapy. Polypeptides and polynucleotides of the invention useful for detecting a cancer in a patient for offering a cancer in a patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1362 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1a;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ovarian tumour polypeptide and prevention and/or treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence represents the ovarian carcinoma GPR39 cDNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention relates
CAGCGACAGGCGGCAGCACAGCACCTGCACGAACACCCCGCCGAAACTGCTGCGAGGACAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AGTTCTCCTTGCAGAGGACTGGCGCGCGGGACGCGAAGAGCGAACGGGCGCTGCACAAAGCG
                                                                                          GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTTGGCCGCAGC
                                                                                                                                                                                                               CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
                                                                                                                                                                                                                                                                                                                                                                                               GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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) STOLK J A.
) ALGATE P A
) FLING S P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   57.4%; ilarity 100.0%; Conservative
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2000US-0656668.
2000US-0713550.
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Pred. No. 2.3e-133;
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RESULT 18
AAD18690/c
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                                                                                                                                           a sample containing a nucleic acid molecule with a hybridisation probe or amplification primers and detecting the presence. The invention is used in gene therapy. The method of the invention is used for identifying nucleic acids or polypeptides associated with a cardiovascular or tumourigenic disorder such as aberrant anglogenesis, aberrant vascularisation, artherosclerosis, or ovarian cancer, ischaemia/ reperfusion injury, hypertension, restenosis, arterial inflammation, endothelial cell disorders, diabetic retinopathy, psoriasis, myocardial infarction, Grave's disease and leukaemia. The methods can also detect mRNA or genomic DNA in a sample. The present sequence is G protein coupled receptor (GPCR) 4941 cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene therapy; aberrant vascularisation; artherosclerosis; ovarian cancer; ischaemia/reperfusion injury; hypertension; arterial inflammation; psoriasis; endothelial cell disorder; diabetic retinopathy; restenosis; myocardial infarction; Grave's disease; G protein coupled receptor; GPCR 4941; vasorropic; hypotensive; antiinflammatory; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CDS
   1256
                                                                                                                                                                                                                                                                                                             The present invention relates to a method for identifying a nucleic actimolecule (G protein coupled receptor gene, GPCR 4941) associated with a cardiovascular or tumourigenic disorder. The method comprising contacti
                                                                                                                                                                                                                                                                                                                                                                                                    cardiovascular and tumorigenic disorders, protein coupled receptor (GPCR)-4941 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-611743/70.
P-PSDB; AAE11751.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human G
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                                                                                                                                                                                                                                                                                                                                                                                                                     Identifying nucleic acids for the diagnosis and treatment of cardiovascular and tumoridenic disorders, comprises identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-APR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antidiabetic; antipsoriatic; leukaemia;
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                                                                                                                                                                                                                                                                                                                                                                          Example 1; Fig 1; 118pp; English.
                                                                                                                    Sequence
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359;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein coupled receptor (GPCR) 4941
                                                                          Similarity
                                                                                                                     2528
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                                                            Conservative
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2000US-0635521.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001WO-US13788
                                                                                                                    BP; 516 A; 766 C; 677 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /product- "Human GPCR 4941" /note- "The CDS is specifically claimed in calim 1 of the specification"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers 42..1403
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                                                                        Score 359;
Pred. No.
                                                            Mismatches
                                                                                                                     567 T; 2 other;
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204 CCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGAGGATCATGTACGCCCGGGAAGT 263

Query Match Best Local Matches

Similarity

56.3%; 99.8%;

Conservative

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Score 352; DB 22; Pred. No. 1.6e-130; 0; Mismatches 1;

DB 22;

Length 444; Indels

0;

Gaps

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RESULT 19
AAH50766/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human tumour
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                                                              patient.
                                                                              are
                                                                                           The
                                                                                                                                           Novel nucleic acid molecules differentially expressed in brain cancers, useful for ascertaining propensity of cell for malignant phenotype or ascertaining suitability of anti-neoplastic drug candidate -
                                                                                                                                                                                                                          Kroes RA,
                                                                                                                                                                                                                                                   (NYXI-) NYXIS NEURO THERAPIES
                                                                                                                                                                                                                                                                             17-NOV-1999;
17-NOV-1999;
                                                                                                                                                                                                                                                                                                                     17-NOV-2000; 2000WO-US31809
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAH50766
Sequence
                           haemangioma, glioblastoma, schwannoma, osteoma and pinealoma. The present sequence is a cancer-associated cDNA of the invention.
                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                181
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                                                present invention provides the sequences of 184 cDNA fragments which differentially expressed in cancer cell depending on the age of the ient. They can be used to diagnose and identify treatments for cers, particularly brain cancers such as haemangibblastoma, teratoma,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
                        sequence is
 444
                                                                                                                    Page 50; 82pp; English.
                                                                                                                                                                                                                           Moskal JR,
BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  associated cDNA #95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                               99US-0166106
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 95
A; 113 C;
                          a cancer-associated
                                                                                                                                                                                                                             Yamamoto H;
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 94 G; 142 T; 0 other;
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RESULT 20
ABL78538/c
ID ABS18/c
ID ABS18/c
ID ABS18/c
AC ABL785
XX ABL785
XX 17-MAY
XX Human;
XX Human;
XX W02001
XX W02001
XX W02001
XX W02001
XX U06-DEC
XX 25-MAY
XX Human;
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The present invention describes a composition (I) comprising: carriers and immunostimulants; and a polypeptide (III) of a ovarian tumour polypeptide encoded by a polypuctode (III) having a cDNA sequence (S1) from the 10912 nucleotide sequences as given in ABL77023 to ABL87934, (III) encoding (II) having a sequence (S2), a T cell population of (II), or antigen presenting cells that express (II). (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to (S1) can be used for detecting ovarian cancer in a patient's biological sample preferably serum or ovarian tissue. The method comprises contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to
                                                                                                                                                                                                                                                                                                                                                                                                                       polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing polypeptide -
                                                                                                                                                                                                                                                                                                                                                                   Claim 1; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Composition for therapy and diagnosis of ovarian cancer comprising
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                                                                                                                                                                                                                                                                                                                                                             ID 1516; 489pp; English.
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                                                    WPI; 2002-122075/16.
                                                                                                                                                                      06-DEC-2001,
                                                                                                                                                                                                                                                                 Human ovarian cancer related cDNA clone SEQ ID NO:2409
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                                                                                                                       26-MAY-2000; 2000US-207484P
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                                                                          Harlocker
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                                                                                                                                                                                                                                                                                                                                                                                                                          606
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100.0%; Pred. No. 2e-115
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polypeptide, polynucleotide encoding to polypeptide or T cell expressing

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RESULT 22
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(I) has cytostatic activity. An oligonucleotide (IV) that hybridises to
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                                             Homo sapiens
                                                                            Human;
                                                                                                                                          17-MAY-2002
                                                                                                                                                                         ABL81262
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA 479
                                                                                                                                                                                                                                                                                                                GACAGTG 606
                                                                                                                                                                                                                                                                                                                                               GACAGTG 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA
                                                                                                                                                                                                                                                                                                                                                                                                           ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              307;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; SEQ ID 2409; 489pp; English
                                                                            ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                          (first entry)
                                                                                                           cancer related cDNA clone SEQ ID NO:4240
                                                                            cancer; ovarian tumour; cytostatic; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         79 A; 84 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           49.18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 307; DB 24;
Pred. No. 1.2e-112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         78 G; 108 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 349;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; SEQ ID 4240; 489pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-122075/16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 409 BP; 89 A; 106 C; 87 G; 127 T; 0 other;
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                                593
15
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                 TGCTTTAGACAGTG
                                                                                  GAACACAGTAAATAAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGC
                                                                                                                                                                                                                                        GAACACAGTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGC
                                                                                                                                                                                                                                                                        ATCAGCCCTGGGCAGACACGAGCAGGAGGGGAGAGACAGAGAAAAAGAAAAACACAGCATGA 412
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                                                                  al Similarity
313; Conserv
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99.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                            Score 263; DB 24;
Pred. No. 3.6e-95;
0; Mismatches 1
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Human ovarian cancer

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CCCNRI-) CORIXA CORP.

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Algate PA, Harlocker

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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      61
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CAATCAGCCCTGGGCAGACACGAGCAGGAGGGGAGACAGAGAAAA
                                                             CAATCAGCCCTGGGCAGACACGAGCAGGAGGAGAGAGAAAA 396
                                                                                                                                                                                                                                                                                             AGAAGGGAAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTCCTTGGGTT 290
                                                                                                                                                                                                                                                           TGGCCGCAGCCATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAA
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                                                                                                                           TGGCCGCAGCCATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAA
                                                                                                                                                                                                                                                                                                                                                                                                    166; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID 8894; 489pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28
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                                                                                                                                                                                                                                                                                                                                                                                                                         Score 166; DB 24; Length 201; Pred. No. 1.4e-56;
                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            46 G;
                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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426 GAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTAC

485 206

GAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTAC

366 AGACACGAGCAGGAGGGAGAGACAGAGAAAAGAAAAACACAGCATGAGAACACAGTAAAT 425

Matches Query Match

Similarity

Conservative (

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Mismatches

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Gaps

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24.6%; Score 154; DB 24; 100.0%; Pred. No. 7.4e-52;

Length 390; Indels

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RESULT 24
ABL78554
                                        CC ABL87934, (III) encoding (II) having a sequence (S2), a T cell (C) population of (II), or antigen presenting cells that express (II). CC (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to CC (S1) can be used for detecting ovarian cancer in a patient's biological contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to can predetermined cutoff value and thereby detecting ovarian cancer in the CC a predetermined cutoff value and thereby detecting ovarian cancer in the CC attent, where the amount of polynucleotide hybridising to (IV) is detected preferably by polymerase chain reaction (PCR). (I) comprising (III) and/or (II) is useful for stimulating and/or expanding T cells competitie for an ovarian tumour protein comprising contacting T cells contacting or (II), (III) is useful in design and preparation of cells and proteins in tumour cells; and to isolate a full length gene from a contacting library e.g., a tumour CDNA library using well known
Sequence
                                                                                                                                                                                                                                                                                                                                             and immunostimulants; and a polypeptide (II) of a ovarian tumour polypeptide encoded by a polynucleotide (III) having a cDNA sequence (S1) from the 10912 nucleotide sequences as given in ABL77023 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CORI-) CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                         present invention describes a composition (I) comprising: carriers
390 BP; 121 A; 82 C;
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99 G; 86 T; 2 other;
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RESULT 25
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Best Local Similarity
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                                                                                                                                                                                                                                       amount of polynucleotide hybridising to (IV) and comparing the amount to a predetermined cutoff value and thereby detecting ovarian cancer in the patient, where the amount of polynucleotide hybridising to (IV) is detected preferably by polynerase chain reaction (PCR). (I) comprising (III) and/or (II) is useful for stimulating and/or expanding T cells specific for an ovarian tumour protein comprising contacting T cells with (III) or (II). (III) is useful in design and preparation of ribozyme molecules for inhibiting expression of the tumour polypeptides and proteins in tumour cells; and to isolate a full length gene from a suitable library e.g., a tumour CDNA library using well known
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing polypeptide -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; ovarian cancer; ovarian tumour; cytostatic; gene;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   (S1) can be used for detecting ovarian cancer in a patient's biological sample preferably serum or ovarian tissue. The method comprises contacting a biological sample from a patient with (IV), detecting the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABL87934, (III) encoding (II) having a sequence (S2), a T cell population of (II), or antigen presenting cells that express (II). (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a composition (I) comprising: carriers and immunostimulants; and a polypeptide (II) of a ovarian tumour polypeptide encoded by a polypucleotide (III) having a cDNA sequence (SI) from the 10912 nucleotide sequences as given in ABL77023 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Algate PA, Harlocker SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-DEC-2001
                                                                                                                                                                                        Sequence 558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID 2375; 489pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CORI-) CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-MAY-2000; 2000US-207484P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-MAY-2001; 2001WO-US17756
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          267
                                                        138 ACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGT 197
 198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CAATTTTTCAGTGTTGGACTTGACAGCTTCTTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002-122075/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CAATTTTTCAGTGTTGGACTTGACAGCTTCTTTT
TGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCC 257
                                      ACAGCACCTGCACGAACACCCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGT 193
                                                                                                              Conservative
                                                                                                                                                                                        BP; 147 A; 133 C; 166 G; 106 T; 6 other;
                                                                                                                                  100.0%;
                                                                                                                                                   19.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jones
                                                                                                                                Score 120; DB 24; pred. No. 2.3e-38;
                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  æ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               519
                                                                                                                                                 Length 558;
                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SS
                                                                                                                0;
                                                                                                                Gaps
                                                                                                                  0
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RESULT 26
ABL84848
                                                                                                 Q
                                    Matches
                                                                 Query Match
                                                                                                                                                                                                ABL87934, (III) encoding (II) having a sequence (SS), a T cell population of (II), or antigen presenting cells that express (II).

(S1) can be used for detecting varian cancer in a patient's biological sample preferably serum or ovarian tissue. The method comprises contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to a predetermined cutoff value and thereby detecting ovarian cancer in the patient, where the amount of polynucleotide hybridising to (IV) is detected preferably by polymerase chain reaction (PCR). (I) comprising (III) and/or (II) is useful for stimulating and/or expanding T cells with (III) or (III) (III) is useful in design and preparation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing polypeptide -
                                                                                                                                         with (III) or (II). (III) is useful in design and processes for inhibiting expression of the tumour polypeptides and proteins in tumour cells; and to isolate a full length gene from a suitable library e.g., a tumour cDNA library using well known
                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a composition (I) comprising: carrier and immunostimulants; and a polypeptide (II) of a ovarian tumour polypeptide encoded by a polynucleotide (III) having a CDNA sequence (SI) from the 10912 nucleotide sequences as given in ABL77023 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; SEQ ID 7826; 489pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human ovarian cancer related cDNA clone SEQ ID NO:7826.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-MAY-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABL84848 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-122075/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Algate PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAY-2000; 2000US-207484P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-MAY-2001; 2001WO-US17756
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                                                                                                   Sequence 381
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CORI-) CORIXA CORP
                                                                                                                                  techniques.
                                                   Local
   ب
   AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCCGCTGCACAAAGCG 60
                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Harlocker
                                  Conservative
                                                                                                     В₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer; ovarian tumour; cytostatic; gene;
                                                                                                     85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cDNA;
                                                                                                   A; 101 C;
                                                 14.7%; Score 92;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       describes a composition (I) comprising: carriers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           381
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jones
                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВP
                                                                                                     122 G;
                                     Mismatches
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                                                                                                     73 T; 0 other;
                                                                    DB 24;
                                                     3.2e-27;
                                       0
                                                                    Length 381;
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GGCGCTGTCGGTGGTGCGCATGTACGC

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GGCGCTGTCGGTGGTGGAGTGCGCATGTACGC

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ID AAL33985
                                                                Matches
                                                                                               Query Match
                                                                                                                                                                                                                                        The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoletin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, croteins, cytochromes, kinesins, cytokines, interferons, interleukins, corporated receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the paptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune
                                                                                                                                                                                        diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polymorphic nucleic acids encoding e.g. amylases, cyclins, poncogenes and histones, useful for diagnosing and treating, antoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer; amyloid protein; anglopoletin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factor; complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                Sequence 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-DEC-1999; 99US-0173419.
27-DEC-2000; 2000US-0173419.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human SNP oligonucleotide #7193.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nervous system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-DEC-2000; 2000WO-US35498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunosuppressive; immunostimulatory; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CURA-) CURAGEN CORP
                                                                                                                                                                             eukaemia),
                             293 GCCGCAGCCATGATCCTCCGAATCTG 318
26
                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2001-465210/50
                                                              26; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 3451; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
                                                                                                                              BP; 14
                                                                                                                                                                             diseases of the nervous system and an infection
                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Leach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disease;
                                                                                                                                A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA; 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                            4.2%;
                                                                                                                              13 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP
                                                                                                                           17 G; 6 T; 0 other;
                                                              0;
                                                                            Score 26;
Pred. No.
                                                              Mismatches
                                                                                 ő
                                                                              . 68;
                                                                                             22;
                                                              0,
                                                                                            Length 50;
                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cyclins, polymerases, treating, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytostatic;
                                                            0,
                                                                                                                                                                          of pathogenic
                                                            Gaps
                                                            0,
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RESULT 28
AAL33986/c
                                                                                                                                          Query Match
Best Local
                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                        proteins, cytochromes, kinesins, cytokines, interferons, interleukins, or-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, the leukaemia), diseases of the nervous system and an infection of pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating fact complement related protein; cytochrome; kinase; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 3451; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAL33986;
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                                                                                                                                                                                                                               Sequence 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoletin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-DEC-1999;
27-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-DEC-2000; 2000WO-US35498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nervous system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human SNP oligonucleotide #7194.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CURA-) CURAGEN CORP.
                                                         267
   50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sapiens.
                                                      ACCTCGTCCAGTCGTGCTTGGGTTTG 292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001-465210/50
   ACCTCGTCCAGTCGTGCTTGGGTTTG
                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RA,
                                                                                                                                                                                                                               BP; 14 A;
                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2000US-0173419
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Leach
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                                                                                                                                          4.2%;
100.0%;
                                                                                                                                                                                                                               12 C; 17 G; 7 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            autoimmune disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP
                                                                                                               0;
                                                                                                                                          Score 26;
Pred. No.
25
                                                                                                               Mismatches
                                                                                                                                          NO.
                                                                                                                                                                                                                            0 other;
                                                                                                                                          DB 22;
0.68;
                                                                                                                                                                   Length 50
                                                                                                               Indels
                                                                                                               0
                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               factor
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AAL27277/c ID AAL27277

standard;

DNA;

51

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RESULT

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ABA18284
ID ABA:
XX
AC ABA:
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                                                                                                                                                                                                                                                                                                         The present invention relates to oligonucleotides encoding polymorphic CC variants of proteins related to amylases, amyloid proteins, angiopoietin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, CC apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, CC fistones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, CC grotein coupled receptors and thicesterases. The present sequence is CC one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of CC diseases associated with inappropriate expression of the proteins listed CC multifactorial diseases with a genetic component, such as autoimmune CC diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, cystemic lupus erythromatosus and Grave's disease), inflammation, cancer CC (e.g. cancers of the bladder, brain, breast, colon and kidney, leakaemia), diseases of the nervous system and an infection of pathogenic
                                                                                  RESULT 30
                                                                                                                                                                                                Query Match
Best Local S
Matches 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thicesterase; inflammation; multifactorial disease; autoimmune disease; infection; nervous system disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neuroprotective; antimicrobial; gene therapy; vaccine; amylase; can amyloid protein; angiopoletin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating f
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and cancer, autoimmune diseases and infections - \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-JUL-2001
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                    ABA18284;
                                                                                                                                                                                                                                                                Sequence 51 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-DEC-1999; 99US-0173419.
27-DEC-2000; 2000US-0173419.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic;
                                                  ABA18284 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CURA-) CURAGEN CORP
                                                                                                                                                 401 AACACAGCATGAGAACACAGTAAAT 425
                                                                                                                                  51
                                                                                                                                   AACACAGCATGAGAACACAGTAAAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001-465210/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP oligonucleotide #485
                                                                                                                                                                                                  25;
                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 1529; 4143pp; English.
                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                    263
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                                                                                                                                                                                                                                Score 25;
                                                                                                                                                                                                                   Pred. No.
                                                                                                                                   27
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                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                T; 0
                                                                                                                                                                                                                                  DΒ
                                                                                                                                                                                                                                                                other;
                                                                                                                                                                                                                                22;
                                                                                                                                                                                                                              Length 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cyclins, polymerases, treating, e.g.
                                                                                                                                                                                                  Indels
                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                of pathogenic
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28-JUN-2000 30-JUN-2000 07-JUL-2000 11-JUL-2000 11-JUL-2000 11-JUL-2000 11-JUL-2000 11-AUG-2000 11-AU

2000US-0214886
2000US-0215135
2000US-0216887
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2000US-0218290
2000US-0224518
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2000US-02231413
2000US-0231441
2000US-0231441
2000US-0231441
2000US-0231441
2000US-0231681
2000US-0231681
2000US-0231681
2000US-0231681

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Human; nootropic; neuroprotective; cytostatic; dermatological; virucide; immunosuppressive; antiinflammatory; anti-HIV; antibacterial; vulnerary; antiparkinsonian; antisickling; antianaemic; antiarthritic; cancer; antirheumatic; hepatotropic; cerebroprotective; antiinflammatory; antiallergic; antidiabetic; antiucer; anticonvulsant; antifungal; antiparasitic; cardiant; immune disorder; cardiovascular disorder; neurological disease; infection; nephrotropic; gene therapy; vaccine; ds
                                                                                                                                                                                                                                                                                                                                                                                       WO200159063-A2
                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-JAN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     system related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polynucleotide
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17-JAN-2001; 2001WO-US01334

2000US-0186350. 2000US-0189874. 2000US-0190076. 2000US-0198123. 2000US-0205515. 2000US-0209467.

2000US-0184664 2000US-0180628

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114-SEP-2000
21-SEP-2000
25-SEP-2000
25-SEP-2000
26-SEP-2000
27-SEP-2000
29-SEP-2000
20-OCT-2000
20-OC
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2000US-0249214
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2000US-0249244
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2000US-0249265
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2000US-0246609

2000US-0246611

2000US-0246613

2000US-0249207

2000US-0249207

2000US-0249209

2000US-0249210

2000US-0249210

2000US-0249211

2000US-0249212
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2000US-0241787.
2000US-0241809.
2000US-0241826.
2000US-0244617.
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2000US-0232400
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Homo sapiens.

07-DEC-2001 AAL10167;

(first entry)

0;

WO200151628-A2

Human; breast cancer; cell marker; cytostatic; ss Human breast cancer expressed polynucleotide 2624.

10-JAN-2001; 2001WO-US00798

19-JUL-2001

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RESULT 31
AAL10167/c
ID AAL10167 standard;
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                                                                                                                              Qy
                                                                                                                                                                                                                                                           The nucleic acids, proteins, antibodies and (ant)agonists are useful?

In the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone, bone cancer, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune chaemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's collease, multiple sclerosis, rheumatoid arthritis and ulcerative collitis; (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing; (e) neurological disease e.g. cerebral anoxia and epilepsy; and (f) infectious diseases such as viral, bacterial, fungal and grarasitic infections.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                  Matches
                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-DEC-2000;

05-DEC-2000;

06-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;

11-DEC-2000;

11-DEC-2000;

15-JAN-2001;
                                                                                                                                                                                                                                     Sequence 263 BP; 88 A; 36 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to novel genes (ABA11004-ABA21534) and proteins (ABB14678-ABB18001) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 10615; 1701pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acids encoding 3224 human nervous system antigen polypeptides, useful for preventing, diagnosing and/or treating nervous system cancers and metastases -
                                                                                                        603 AGTGNAAAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-541565/60.
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                                                                                           241 AGTGNAAAAAAAAAAAAAAAA 263
                                                                                                                                                                                   Local Similarity
                                                                                                                                                                    23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Barash SC,
                                                                                                                                                              3.7%; Score 23; llarity 100.0%; Pred. No. Conservative 0; Mismatc
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2000US-0251479.

2000US-0251868.

2000US-0251868.

2000US-0251869.

2000US-0251869.

2000US-0251990.

2000US-0251990.

2000US-0254097.

2001US-0259678.
   cDNA; 377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ruben SM;
                                                                                                                                                                                                                                   67 G;
                                                                                                                                                              red. No. 8.
Mismatches
                                                                                                                                                                                                                                   71 T; 1 other;
                                                                                                                                                                             DB 22; Length 263; 8.7;
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RESULT 32
ABV23163
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14-MAR-2000; 2000US-0189167.
24-MAR-2000; 2000US-0193099.
29-MAR-2000; 2000US-0193480.
15-MAY-2000; 2000US-0205230.
09-JUN-2000; 2000US-0211315.
25-JUL-2000; 2000US-0220534.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to human breast cancer expressed polynucleotides (AALO7544-AAL26789) and methods of assessing whether a patient is afflicted with breast cancer by examining the correlation between the expression of certain markers and the cancerous state of breast cells. The polynucleotides and encoded polypeptides are potential markers for detecting, diagnosing, monitoring, characterising treating and potentially preventing breast cancer. The polynucleotides and encoded polypeptides are also useful for isolating compounds with cytostatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 496;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lillie J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 377 BP; 87 A; 62 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New peptide useful as a marker for the diagnosis of breast cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                                                                                                                      23-AUG-2001.
                                                                                                                                                                                                              WO200160860-A2
                                                                                                                                                                                                                                                         pharmacogenomic
                                                                                                                                                                                                                                                                                            Human prostate expression marker cDNA 23154.
                                                                                                                                                                                                                                                                                                                    16-SEP-2002
                                                                                                                                                                                                                                                                                                                                            ABV23163;
                                                                                                                                                                                                                                                                                                                                                                  ABV23163 standard; cDNA; 1539
                                                                                           09-JUN-2000;
18-JUL-2000;
                                                                                                                                         17-FEB-2000;
                                                                                                                                                                20-FEB-2001;
                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                        Human;
                                 Schlegel R,
                                                       (MILL-)
                                                                                                                 25-MAY-2000;
                                                                                                                              16-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                            107
                                                                                                                                                                                                                                                                                                                                                                                                                                                  603 AGTGNAAAAAAAAAAAAAAAAA 625
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           2001-662795/76
                                                                                                                                                                                                                                                                    prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                        MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Xu Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                             2000US-183319P.
2000US-189862P.
2000US-207454P.
2000US-211314P.
2000US-211314P.
2000US-255281P.
                                                                                                                                                                                                                                                                                                                    (first entry)
                                 Endege
                                                                                                                                                                2001WO-US05171
                                                                                                                                                                                                                                                         cancer; cytostatic; carcinogen; pharmacodyanamic marker;
marker; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3695pp; English.
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M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.7%;
                                  Monahan JE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Steinmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        87 G; 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. 8. Mismatches
                                                                                                                                                                                                                                                                                                                                                                   ВР
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 22; Length 377;
8.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25 other;
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Best Local
                                                                                                                                                                                    17-FEB-2000; 2000US-183319P.
16-MAR-2000; 2000US-189862P.
25-MAY-2000; 2000US-207454P.
09-JUN-2000; 2000US-211314P.
18-JUL-2000; 2000US-219007P.
13-DEC-2000; 2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:
(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usel for detecting presence of prostate cancer, stage of prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 4154; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1539 BP; 577 A; 372 C; 435 G; 146 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (e) selecting a composition for inhibiting prostate cancer in a patien (f) assessing the prostate cell carcinogenic potential of a compound;(g) determining whether prostate cancer has metastasized in a patient;(h) assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (d) assessing the efficacy of a therapy for inhibiting prostate cancer
in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                       pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human prostate expression marker cDNA 28994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABV29003;
                                                                                                                                                                                                                                                                                                                     23-AUG-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABV29003 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      patient;
                                       Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usel for detecting presence of prostate cancer, stage of prostate cancer
                                                                                                                                                                                                                                                                                          20-FEB-2001; 2001WO-US05171
                                                                                                  WPI; 2001-662795/76.
                                                                                                                               Schlegel R,
                                                                                                                                                         (MILL-) MILLENNIUM PREDICTIVE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                878
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             385 AGACAGAGAAAAGAAAAACACAG 407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         is also useful as a pharmacodyanamic or pharmacogenomic marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGACAGAGAAAAGAAAAACACAG
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                                                                                                                                                                                                                                                                                                                                                                                                                       prostate
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                                                                                                                                 Endege
                                                                                                                                                                                                                                                                                                                                                                                                                          cancer;
                                                                                                                                                                                                                                                                                                                                                                                                            marker; gene;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                          cytostatic; carcinogen; pharmacodyanamic marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1539
                                                                                                                                 Monahan JE;
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Pred. No.
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7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                prostate cancer in a patient;
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useful
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useful
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Claim

1; Page 6134; 11750pp; English

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RESULT 34
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
     The invention relates to an isolated polynucleotide (I) comprising sequences selected from 273 sequences fully defined in the specification (I), including its encoded polypeptide (II), an antibody binding to (II), a fusion protein comprising (II) and a T-cell population stimulated by (I) or (II) are useful for stimulating an immune response in a patient and treating head and neck cancer in a patient. An oligonucleotide (III) that hybridises to (I) is useful for determining the presence of cancer in a patient, by obtaining a biological sample from the patient, contacting the sample with (III), detecting in the sample an amount of a polynucleotide that hybridises to the oligonucleotide, and comparing the
                                                                                                                                                                                                                                                          Wang
                                                                                                                                                      Claim 1; Page
                                                                                                                                                                                                         Novel polynucleotide
                                                                                                                                                                                                                                  WPI; 2002-257467/30
                                                                                                                                                                                                                                                                                                               03-AUG-2000;
16-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                    01-AUG-2001; 2001WO-US24226
                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer in a patient;
(d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
(e) selecting a composition for inhibiting prostate cancer in a patien;
(f) assessing the prostate cell carcinogenic potential of a compound;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:
(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                     (CORI-)
                                                                                                                                                                                                                                                                                                                                                                                14-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human head and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK53839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABK53839 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                878
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         34
                                                                                                                                                                                          pharmaceutical compositions, e.ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              determining whether prostate cancer has metastasized in a patient; assessing the aggressiveness or indolence of prostate cancer is tent;
                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGACAGAGAAAAGAAAAACACAG 900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AGACAGAGAAAAGAAAAACACAG 407
                                                                                                                                                                                                                                                                                                                                                                                                                                                             SS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           head
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          also useful as a pharmacodyanamic or pharmacogenomic marker
                                                                                                                                                                                                                                                                                     CORIXA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                          Fan L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                               2000US-223281P
2000US-249933P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP;
                                                                                                                                                      123;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    neck
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neck
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    577 A;
                                                                                                                                                    200pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tumour cDNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    372 C;
                                                                                                                                                     English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 23;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tumour;
                                                                                                                                                                                          e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID No
                                                                                                                                                                                          and neck tumour
g. vaccines, for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <u>ი</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 1539;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immunogenic;
                                                                                                                                                                                       polypeptides, treating head
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          patient;
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t of the
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RESULT 35
ABV08332/c
ID ABV08
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Best Local S
Matches 22
                                 cancer in a pat
(d) assessing t
in a patient;
(e) selecting a
(f) assessing t
(g) determining
(h) assessing t
                                                                                        The invention relates to an isolated nucleic acid molecule (I) com a nuclectide sequence given in Tables 1-9 (ABV00010-ABV62213) of t specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient;

(a) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
                                                                                                                                                                                                      Claim 1;
                                                                                                                                                                                                                         Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usef for detecting presence of prostate cancer, stage of prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amount of polynucleotide that hybridises to the oligonucleotides to a predetermined cut-off value. (I) and (II) are useful in pharmaceutical compositions, e.g. vaccines, and other compositions for the diagnosis; treatment of head and neck cancer. ABK53806-ABK54078 represent human head and neck cancer of the invention.
                                                                                                                                                                                                                                                                               WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                  17-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABV08332 standard; cDNA; 377
                                                                                                                                                                                                                                                                                                                                                                  09-JUN-2000;
18-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                          16-MAR-2000;
25-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200160860-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABV08332;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                             (MILL-) MILLENNIUM PREDICTIVE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                         20-FEB-2001; 2001WO-US05171
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                                                                                                                                                                                                                                                                                                      ₽,
                                                                                                                                                                                                    Page 1324; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               256 BP; 83 A; 31 C;
                                                                                                                                                                                                                                                                                                                                                    ; 2000US-183319P.
2000US-189862P.
2000US-207454P.
2000US-211314P.
2000US-219007P.
2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                    Endege
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cytostatic; carcinogen; pharmacodyanamic marker;
                                                                                                                                                                                                                                                                                                      Monahan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            marker cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        625
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                    JE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22;
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22;
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                                                                                                                                                                  ) comprising of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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selecting a composition for inhibiting prostate cancer i assessing the prostate cell carcinogenic potential of a determining whether prostate cancer has metastasted in assessing the aggressiveness or indolence of prostate cancers as the cancer has metastasted in assessing the aggressiveness or indolence of prostate cancer has metastasted.

prostate cancer

'n

a patient;

useful

as

pharmacodyanamic or

pharmacogenomic marker

cancer in

a patient;

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226 GTGNAAAAAAAAAAAAAAAAA 247

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RESULT 36
AAI87949
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                                                      Query Match
Best Local
                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 377 BP; 89 A; 75 C;
                                                                                                                                                                                     The invention relates to human polynucleotides (AAI79941-AAI93841) and the encoded proteins (AA000010-AA013910) that exhibit activity elating to cytokine, cell proliferation or cell differentiation or which may induce production of other cytokines in other cell populations. The polynucleotides and polypeptides are useful in gene therapy, vaccines or peptide therapy. The polypeptides have various cytokine-like activities, e.g. stem cell growth factor activity, haematopoisels regulating activity, tissue growth factor activity, immunomodulatory activity and activinyinhibin activity and may be useful in the diagnosis and/or representation of cancer lawysement acresses exercise diagnosis and/or representations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; cytokine; cell proliferation; cell differentiation; gene th vaccine; peptide therapy; stem cell growth factor; haematopoiesis; tissue growth factor; immunomodulatory; cancer; leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tissue growth factor; immunomodulatory; cancer; leukaemia; nervous system disorders; arthritis; inflammation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAI87949 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-FEB-2000;
18-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-SEP-2001
                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID NO 8009; 1399pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                          disorders
                                                                                                                                                                                                                                                                                                                                                                      Isolated nucleic acids and polypeptides, useful for preventing diagnosing and treating e.g. leukaemia, inflammation and immune
                                                                                                                                                                                                                                                                                                                                                                                                               P-PSDB; AA008018.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HYSE-) HYSEQ INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-FEB-2001; 2001WO-US04927.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200164835-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                    Note: The sequence data for this patent did not form specification, but was obtained in electronic format
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          604 GTGNAAAAAAAAAAAAAAAAA 625
                                                                                                                                                                               treatment of cancer,
                                                                                              Sequence
                                                                                                                                                                  inflammation.
              604 GTGNAAAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  78
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                                                                                                                         ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GTGNAAAAAAAAAAAAAAAAA 57
                                                                                                                                                                                                                                                                                                                                                                                                                             2001-514838/56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polynucleotide SEQ ID NO 8009
Similarity
                                                      Similarity
                                                                                               442
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Liu C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000US-0515126.
2000US-0577409.
                                                                                              BP;
                                                                                               167
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                                                      3.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                                                                                                                                                                               leukaemia,
                                                                                               Α;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.58;
                                                                                               65 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 22;
Pred. No.
                                                      Score 22;
Pred. No.
                 625
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                                                                                               102 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                               nervous system disorders, arthritis and
                                          mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cell differentiation; gene therapy;
                                                                                                85 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 23; Length 377; 21;
                                                                     DB
                                                         20;
                                                                     22;
                                                                                                  23 other;
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                                            0;
                                                                     Length 442;
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                                            Indels
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                                Qy
                                                                   Matches
                                                                                                    Query Match
                                                                                                                                                                                                                                    The nucleic acids, proteins, antibodies and (antiagonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, rheumatoid arthritis and ulcerative colitis, (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and epilepsy; and (f) infectious diseases such as viral, bacterial, fungal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral; antiallergic; hepatotropic; antidiabetic; antiinflammatory; antiulcer; vulnerary; anticonvulsant; antibacterial; antifungal; antiparasitic; cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABL90482 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel 1405 isolated polypeptides, useful for diagnosis, treatment and prevention of neural, immune system, muscular, reproductive, gastrointestinal, pulmonary, cardiovascular, renal and proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                neurological disease; infection; human; secreted protein; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human polynucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                            (ABB89040-ABB90444) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Birse CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-MAY-2000; 2000US-205515P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                      Sequence 1329 BP;
                                                                                                                                                                    Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format direction with at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to novel genes (ABL89449-ABL90853) and proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; SEQ ID NO 1044; 2081pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-122018/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HUMA-) HUMAN GENOME
                                604 GTGNAAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                                                        parasitic infections.
ABB90073
                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rosen CA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CDNA; 1329
                                                                                                                                       263 A; 407 C; 369 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SCI INC
                                                                                     100.0%;
                                                                                                         3.5%;
                                                                     Score 22; DB
s; Pred. No. 18;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NO 1044.
   1315
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                                                                                                                                           288
                                                                                                         DB
                                                                                                                                                                                           in electronic format directly
                                                                                         18;
                                                                                                         24;
                                                                                                                                           T; 2 other;
                                                                       0;
                                                                                                       Length 1329;
                                                                         Indels
                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        disorder;
                                                                         Gaps
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cc encoding an Arabidopsis thaliana transcription factor, their variants, complements, fragments, or related polynucleotide with 31% to 95% cc sequence identity, where the plant possesses an altered trait as compared to a wild-type or reference plant, or the plant axhibits an altered cc phenotype as compared to a wild-type or reference plant, or the plant compared to a wild-type or reference plant, or the plant or associated with a plant trait as compared to a wild plant. Also included created with a plant comprising the polynucleotides, a computer cc associated with a plant trait as compared to a wild plant. Also included created with a plant trait as compared to a wild plant. Also included created with a plant trait as compared to a wild plant. Also included created with a plant trait as compared to a wild plant. Also included created with a plant trait as compared to a wild plant. Also included created by a plant trait as compared to a wild plant. Also included creating a plant trait sequence information, and identifying a composition of plant trait sequence information, and identifying a created by plant and identifying a plant to plant sequence of compissing a plant plant and identifying a polynucleotide or an antisense nucleic acid, inserting the polynucleotide or antisense nucleic acid, inserting the compission the vector, and selecting for a modified trait (e:g. increased production of agriculturally useful proteins or metabolic chemicals,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 38
ABK65378
                                                                                                                                                                                                                                                                                                                             The invention relates to 1 of 232 isolated or recombinant polynucleotides encoding an Arabidopsis thaliana transcription factor, their variants,
                                                                                                                                                                                                                                                                                                                                                                              Claim 4; Page 927-929; 941pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                            An isolated or recombinant polynucleotide used to produce a transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pilgrim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-AUG-2000; 2000US-227439P.
16-NOV-2000; 2000US-0713994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200215675-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABK65378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (PINE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-APR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-AUG-2001; 2001WO-US26189.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fruit yield; growth rate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             agriculture;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Plant; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Arabidopsis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABK65378 standard; cDNA; 1388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RIEC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RATC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ADAM/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HEAR/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CREE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (PILG/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REUB/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     griculture; metabolic chemical; environmental stress; drought;
icrobial disease resistance; herbicide resistance; seed yield;
ruit yield; growth rate; leaf senescence; flower senescence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PSDB; AAU93192.
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CREELMAN R.
DUBELL A J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RATCLIFF O. REUBER J L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HEARD J.
JIANG C.
KEDDIE J.
ADAM L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    M, Creelman
Ratcliff O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PINEDA O.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MENDEL BIOTECHNOLOGY INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RIECHMANN J L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene; transcription factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cDNA encoding a transcription factor #230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Reuber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dubell AJ, Hear JL, Riechmann JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        transgenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jiang C,
L, Yu G,
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Pineda O;
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RESULT 3
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                                                                           expression. For example, N and P may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of P by expressing inactive proteins or to supplement the patients own production of P. Additionally, N may be used to produce the colon cancer-associated Ps, by inserting the nucleic acids into a host cell and culturing the cell to express the proteins. N and P can be used in the prevention, diagnosis and treatment of colorectal carcinomas and cancers. AAH37196 to AAH37204 and AAB77789 represent sequences used in the exemplification of the
     present invention.

N.B. Pages 666 to 682 and page 7053 of the sequence listing were missing at time of publication, meaning no sequences are present SEQ ID NO:1027 to 1052, 7921 and 7922.
                                                                                                                                                                                                                                    cancer-associated nucleic acid molecules (N) and proteins (P), where the proteins are collectively known as to concer antigens. The colon cancer antigens have cytostatic activity and can be used in gene therapy and vaccine production. N and P may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate P
                                                                                                                                                                                                                                                                                                                           AAH32943 to AAH37195 and AAG73514 to AAG77788 represent human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       pest tolerance, environmental stress response (e.g. drought), microbial disease resistance, herbicide resistance, seed and fruit yield, growth rate, leaf and flower senescence and many other traits listed in the specification). The present sequence is one of the 232 polynucleotides encoding an A. thaliana transcription factor.
                                                                                                                                                                                                                                                                                                            cancer-associated nucleic acid molecules (N)
                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acids encoding 4277 human colon cancer-associated polypeptides, useful for preventing, diagnosing and/or treating colorectal cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                       P-PSDB; AAG74876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-235357/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ruben SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-SEP-1999;
03-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     colorectal carcinoma; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1388 BP; 398 A; 282 C; 319 G; 389 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; colon cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH34281;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
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                                                                                                                                                                                                                                                                                                                                                             Page 3077-3078; 9803pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-0157137.
99US-0163280.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  colon cancer antigen; diagnosis; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Birse CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     encoding cDNA SEQ ID NO:1363.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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RESULT 40
ABL90818/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral; antiallergic; hepatotropic; antidiabetic; antiinflammatory; antifucer; vulnerary; anticonvulsant; antibacterial; antifungal; antibarastic; cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder; neurological disease; infection; human; secreted protein; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1823 BP; 355 A; 609 C;
                                                                                                                                                              (ABB89040-ABB90444) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune hademolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-MAY-2001; 2001WO-US16450.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABL90818 standard;
                                                                             (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and epilepsy; and (f) infectious diseases such as viral, bacterial, funga and parasitic infections.
                                                                                                                                                                                                                                                                                                                                                                                               Novel 1405 isolated polypeptides, useful for diagnosis, treatment and prevention of neural, immune system, muscular, reproductive, gastrointestinal, pulmonary, cardiovascular, renal and proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-MAY-2000; 2000US-205515P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200190304-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                   disease, multiple sclerosis, rheumatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as myocardial ischaemias;
                                                                                                                                                                                                                                                                                                                                                    Claim 4; SEQ ID NO 1380; 2081pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HUMA-) HUMAN GENOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 604 GTGNAAAAAAAAAAAAAAAA 625
 Sequence
                                                   printed specification, but was
                                from WIPO
                                                                                                                                                                                                                                                                                                                     invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GTGNAAAAAAAAAAAAAAAAA 1747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2002-122018/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polynucleotide SEQ ID NO 1380.
                                                                 The sequence data
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABB90409.
                                at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rosen
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                                                                                                                                                                                                                                                                                                                      relates
   BP;
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   383 A; 571 C; 686 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cDNA;
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                                                                                                                                                                                                                                                                                                                   to novel genes (ABL89449-ABL90853) and proteins
                                                  for this patent did not form part of but was obtained in electronic format
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2049
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17;
   400 T; 9 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22;
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                                                   part of the format directly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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                                                                                                     fungal
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Query Match

22;

DB

24;

Length 2049;

07-DEC-2001

(first entry)

AAL09957; AAL09957

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standard;

CDNA; 288

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RESULT 41
AAH69474/c
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RESULT 42
AAL09957/c
ID AAL09
XX
AC AAL09
XX
DT 07-DE
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                                                                                                                                            Matches
                                                                                                                                                       Query Match
Best Local :
                                                                                                                                                                                                        The invention relates to novel genes (AAH68727-AAH73383) associated with cervical cancer with cytostatic activity. The nucleic actis and encoded polypeptides are useful: to assess if a patient is afflicted with cervical cancer or has a pre-malignant condition; to monitor the progression of cervical cancer or a premalignant condition in a patient; and to select and/or assess the efficacy of a compound or therapy for inhibiting cervical cancer in a patient. The nucleic acids may also be useful for gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH69474 standard; cDNA; 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human cervical cancer marker nucleic acid 748
                                                                                                                                                                                                                                                                                                                                   and
                                                                                                                                                                                                                                                                                                                                              New
                                                                                                                                                                                                                                                                                                                                                                                                                                  09-JUN-2000;
21-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-MAR-2000;
12-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-DEC-1999;
21-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-JUN-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200142467-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cervical cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-SEP-2001
                                                                                                                                                                                        Sequence 259 BP; 73 A; 33 C;
                                                                                                                                                                                                                                                                                                             Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                             (MILL-) MILLENNIUM PREDICTIVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-DEC-2000;
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                                                                                                                      605 TGNAAAAAAAAAAAAAAAAA 625
                                                                                                  63
                                                                                                                                                                                                                                                                                                                                  isolated nucleic acid for diagnosing and treating cervical cancer for assessing and detecting compounds for treating the cancer \cdot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GTGNAAAAAAAAAAAAAAAA 304
                                                                                                                                                                                                                                                                                                                                                                    2001-375006/39
                                                                                                   TGNAAAAAAAAAAAAAAAA 43
                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                         ₩,
                                                                                                                                                                                                                                                                                                             Page 232; 1051pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000US-0189315.
2000US-0203791.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000WO-US33312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                         Deeds J,
                                                                                                                                                                                                                                                                                                                                                                                                                                    2000US-0210600.
2000US-0220114.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.08; ...
                                                                                                                                                        100.0%;
                                                                                                                                                                   3.48;
                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pre-malignant condition;
                                                                                                                                                        Score 21;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
                                                                                                                                                                                         37 G;
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                                                                                                                                                         54;
                                                                                                                                                                   DB 22;
                                                                                                                                                                                          29
                                                                                                                                                                                          other;
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                                                                                                                                                0,.
                                                                                                                                                                   Length 259;
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                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                        RESULT 43
                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-JAN-2000; 2000US-0176077.
14-MAR-2000; 2000US-0189167.
24-MAR-2000; 2000US-0193099.
29-MAR-2000; 2000US-0193480.
15-MAY-2000; 2000US-0205230.
09-JUN-20000; 2000US-020534.
          20-FEB-2001; 2001WO-US05171.
                                                        WO200160860-A2
                                                                               Homo sapiens
                                                                                                    Human; prostate cancer; cytostatic; carcinogen; pharmacogenomic marker; gene; ss.
                                                                                                                                      Human prostate expression marker cDNA 5082.
                                                                                                                                                                                                            ABV05091 standard;
                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to human breast cancer expressed polynucleotides (AALO7544-AAL26789) and methods of assessing whether a patient is afflicted with breast cancer by examining the correlation between the expression of certain markers and the cancerous state of breast cells. The polynucleotides and encoded polypeptides are potential markers for detecting, diagnosing, monitoring, characterising treating and potentially preventing breast cancer. The polynucleotides and encoded polypeptides are also useful for isolating compounds with cytostatic
                                23-AUG-2001.
                                                                                                                                                                 13-SEP-2002
                                                                                                                                                                                         ABV05091;
                                                                                                                                                                                                                                                                                                                                                                       Sequence 288 BP; 76 A; 53 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 462; 3695pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New peptide useful as a marker for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lillie J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-JAN-2001; 2001WO-US00798
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human breast cancer expressed polynucleotide 2414.
                                                                                                                                                                                                                                                                                              605 TGNAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                           70
                                                                                                                                                                                                                                                                                                                                    Local Similarity
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                                                                                                                                                                                                                                                                        TGNAAAAAAAAAAAAAAA 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001-451856/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                breast cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Xu Y,
                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wang Y,
                                                                                                                                                                                                             cDNA; 288
                                                                                                                                                                                                                                                                                                                                3.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cell marker; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Steinmann
                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                    75 G;
                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                 Score 21;
Pred. No.
                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                    79 T; 5 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diagnosis of breast cancer
                                                                                                                                                                                                                                                                                                                                 DB 22; Length 288; 54;
                                                                                                                                                                                                                                                                                                                     0,
                                                                                                            pharmacodyanamic marker;
                                                                                                                                                                                                                                                                                                                     Indels
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ABV05770/c
ID ABV057
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local :
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09-JUN-2000;
18-JUL-2000;
13-DEC-2000;
                                      17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
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16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                            20-FEB-2001;
                                                                                                                    23-AUG-2001;
                                                                                                                                                WO200160860-A2
                                                                                                                                                                                                   pharmacogenomic
                                                                                                                                                                                                             Human; prostate cancer; cytostatic; carcinogen;
                                                                                                                                                                                                                                       Human prostate expression marker cDNA 5761.
                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                     13-SEP-2002
                                                                                                                                                                                                                                                                                                 ABV05770;
                                                                                                                                                                                                                                                                                                                        ABV05770 standard; cDNA; 294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 288 BP; 76 A; 37 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, use for detecting presence of prostate cancer, stage of prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 864; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Schlegel R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (d) assessing the efficacy of a therapy for inhibiting prostate cancer
in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                    605 TGNAAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                             55 TGNAAAAAAAAAAAAAAA 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  selecting a composition for inhibiting prostate cancer in a patient; assessing the prostate cell carcinogenic potential of a compound; determining whether prostate cancer has metastasized in a patient; assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          is also useful as a pharmacodyanamic or pharmacogenomic marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001-662795/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                        2000US-183319P.
2000US-189862P.
2000US-207454P.
2000US-211314P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; 2000US-183319P.
; 2000US-189862P.
; 2000US-207454P.
; 2000US-211314P.
; 2000US-219007P.
; 2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                          2001WO-US05171
                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                   marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; E
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                                                                                                                                                                                                   gene;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                         No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 23; Length 288; 54;
                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>,,</u>
                                                                                                                                                                                                            pharmacodyanamic marker;
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AAH68974/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:
(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 962; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful for detecting presence of prostate cancer stage of prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Schlegel R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 294 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (e) selecting a composition for inhibiting prostate cancer in a patient;(f) assessing the prostate cell carcinogenic potential of a compound;(g) determining whether prostate cancer has metastasized in a patient;(h) assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (d) assessing the efficacy of a therapy for inhibiting prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                             AAH68974 standard; cDNA; 359 BP
                                                                                                                                                                                         08-DEC-2000;
                                                                                                                                                                                                                                             WO200142467-A2.
                                                                                                                                                                                                                                                                                                   Cervical cancer;
                                                                                                                                                                                                                                                                                                                           Human cervical cancer marker nucleic acid 248.
                                                                                                                                                                                                                                                                                                                                                         19-SEP-2001
              WPI; 2001-375006/39
                                       Schlegel R,
                                                                                                                      14-MAR-2000;
12-MAY-2000;
                                                                                                                                                              08-DEC-1999;
                                                                                                                                                                                                                    14-JUN-2001.
                                                                 (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
                                                                                           21-JUL-2000;
                                                                                                                                                 21-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
les 21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                is also useful as a pharmacodyanamic or pharmacogenomic marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             patient
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                           2000US-0189315.
2000US-0203791.
2000US-0210600.
2000US-0220114.
                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                         2000WO-US33312
                                       Deeds J,
                                                                                                                                                 99US-0169681.
99US-0171350.
                                                                                                                                                                                                                                                                                                   cytostatic; pre-malignant condition; gene therapy; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         79 A; 52 C; 34 G; 119 T; 10 other;
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                                       Berger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Monahan JE;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                          Zhao
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53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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Search completed: November Job time: 142.06 secs
                                                                                                                                               Matches
                                                                                                                                                                  Best
                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated nucleic acid for diagnosing and treating cervical cancer and for assessing and detecting compounds for treating the cancer \cdot
                                                                                                                                                                                                                 Sequence 359 BP; 96 A; 62 C; 48 G;
                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 147; 1051pp; English.
                                                                                                         605 TGNAAAAAAAAAAAAAAAAA 625
                                                                         82 TGNAAAAAAAAAAAAAAAAAA 62
                                                                                                                                                                Local
                                                                                                                                                                  Similarity 100.0%;
                                                                                                                                                 Conservative
                                                                                                                                                                                       3.48;
                7,
                      2002,
                                                                                                                                                   0
                                                                                                                                                                  Score 21; DB 22;
Pred. No. 52;
                        18:15:34
                                                                                                                                                   Mismatches
                                                                                                                                                                                                                          131 T; 22 other;
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Result
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Word size :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Scoring table:
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      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            seq length: 0 seq length: 2000000000
      US-09-970-966-210
625
1 agttctccttgcagag
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Gapop 60.0 , Gapext 60.0
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                                                                                                                                                                                                                                                                                                                                                                                                              /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/pcTUS_COMB.seq:*
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| h<br>Simi<br>21;                      | CURRENT APPLICATION DATA: APPLICATION NUMBER: US/V FILING DATE: 07-OCT-199: CLASSIFICATION: 424 ATTORNEY/AGENT INFORMATION NAME: Maki, David J. REGISTRATION UNMBER: 31, REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMAT: TELEPHONE: (206) 682-6031 TELEPHONE: (206) 682-6031 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 377 base pairs TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: Linear TOPOLOGY: Linear                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  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RESULT 2 US-09-385-982-519/c

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RESULT 3
US-09-149-476-67
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APPLICANT: Rosen et
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TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
TITLE OF INVENTION: PRODUCTS: II
FILE REFERENCE: CCDNA-280XX
CURRENT APPLICATION NUMBER: US/09/385,982
CURRENT FILING DATE: 1999-08-30
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APPLICATION NUMBER: 60/056,884
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FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/ APPLICATION NUMBER:

60/043,674 60/043,671

APPLICATION NU FILING DATE: 1

NUMBER:

60/043,311

997-04-11

APPLICATION NUMBER: 60/043,569 FILING DATE: 1997-04-11

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NUMBER: 6

60/043,314 60/043,568

DATE:

1997-04-

FILING DATE: 1997-04-APPLICATION NUMBER: 60 FILING DATE: 1997-04-APPLICATION NUMBER: 60

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NUMBER: 60

60/043,312 60/043,669

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NUMBER: 60/1: 1997-06-06

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60/043,315

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NUMBER: 60/043,672

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60/056,882 60/056,872 60/056,662 60/056,878 60/056,630 60/056,893 60/056,889 60/056,877 60/056,886 60/048,974

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GENERAL INFORMATION:
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Best Local :
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APPLICATION NUMBER: US 08/844,419

FILING DATE: 18-APR-1997

PRIOR APPLICATION UMBER: US 08/846,017

FILING DATE: 25-APR-1997

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

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MEDIJIM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/974,549A
FILING DATE: 19-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
                                                                 FILING DATE: 06-MAY-1997
PRIOR APPLICATION DATA:
                PRIOR APPLICATION DATA:
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CITY: San Francisco
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                               APPLICATION NUMBER: US 0 FILING DATE: 09-MAY-1997
                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
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R APPLICATION NUMBER: 60/057,669

ER FILING DATE: 1997-09-05

ER APPLICATION NUMBER: 60/049,610

ER FILING DATE: 1997-06-13

ER APPLICATION NUMBER: 60/061,060

ER FILING DATE: 1997-10-02
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Harley, Calvin B.
Andrews, William H.
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Nakamura, Toru
Chapman, Karen B.
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NUMBER:
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                                                  US 08/854,050
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US 08/911,312
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Pred. No.
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US-08-974-549A-3
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                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Rosen et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 309, Application US/09149476 Patent No. 6420526
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Best Local Similarity 100.
Matches 21; Conservative
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REFERENCE/DOCKET NUMBER: 01:
TELECOMMUNICATION INFORMATION:
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LOCATION: 1..2176
OTHER INFORMATION:
FEATURE:
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APPLICATION NUMBER:
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TOPOLOGY: 11
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            ER APPLICATION NUMBER: 60/040,162
ER FILING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/040,333
ER FILING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/038,621
ER FILING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/040,626
ER FILING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/040,334
ER FILING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/040,336
ER FILING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/040,336
ER FILING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/040,163
ER FILING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/040,163
ER FILING DATE: 1997-03-07
ER APPLICATION NUMBER: 60/040,163
ER FILING DATE: 1997-03-07
                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/US98/04493
FILING DATE: 1998-03-06
APPLICATION NUMBER: 60/047,600
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100.0%; Pred. No. 1.5;
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ER APPLICATION NUMBER: 60/043,311
ER FILLING DATE: 1997-04-11
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ER APPLICATION NUMBER: 60/043,674
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ER FILLING DATE: 1997-06-06
ER APPLICATION NUMBER: 60/056,886
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R FILING DATE: 1997-05-23
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R APPLICATION NUMBER: 60/047,631
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APPLICATION NUMBER: 60/043,580
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R APPLICATION NUMBER: 60/0
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ER APPLICATION NUMBER: 60/047,588
ER FILLING DATE: 1997-05-23
ER APPLICATION NUMBER: 60/047,585
ER FILLING DATE: 1997-05-23
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ER FILLING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,845
ER FILLING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/056,892
ER FILLING DATE: 1997-08-22
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ER APPLICATION NUMBER: 60/057,761
ER FILLING DATE: 1997-08-22
ER APPLICATION NUMBER: 60/047,595
ER FILLING DATE: 1997-05-23
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IR APPLICATION NUMBER: 60/05
IR FILING DATE: 1997-08-22
IR APPLICATION NUMBER: 60/05
IR FILING DATE: 1997-08-22 PR APPLICATION NUMBER: 60/047,501
FR FILING DATE: 1997-05-23
FR APPLICATION NUMBER: 60/043,670
FR FILING DATE: 1997-04-11
FR APPLICATION NUMBER: 60/056,632
FR FILING DATE: 1997-08-22 PR FILING DATE: 1997-08-22
FR APPLICATION NUMBER: 60/056,874
FR FILING DATE: 1997-08-22
FR APPLICATION NUMBER: 60/056,910
FR FILING DATE: 1997-08-22
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FR FILING DATE: 1997-08-22 FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,903
APPLICATION NUMBER: 60/056,808 APPLICATION NUMBER: 60/ FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/ FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/1 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/1 FILING DATE: 1997-05-23 FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,911
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,636 APPLICATION NUMBER: 60/056,880 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,894 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,872 FILING DATE: 1997-08-22 NUMBER: 60/047,589: 1997-05-23 1997-05-23 60/043,576 60/043,578 60/047,593 60/047,614 60/056,879 60/056,888 60/056,637 60/056,882 60/056,662 60/056,893

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EARLIER FILING DATE: 1997-08-22
EARLIER APPLICATION NUMBER: 60/056,875
EARLIER FILING DATE: 1997-08-22
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EARLIER APPLICATION NUMBER: 60/056,887
EARLIER FILING DATE: 1997-08-22
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EARLIER APPLICATION NUMBER: 60/056,908
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Best Local :
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EARLIER FILING DATE: 1997-09-05
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EARLIER APPLICATION NUMBER: 60/056,876
EARLIER FILING DATE: 1997-08-22
EARLIER APPLICATION NUMBER: 60/056,881
EARLIER FILING DATE: 1997-08-22
                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/686,878A
FILING DATE:
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
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APPLICANT: Treacy, Maurice
APPLICANT: Evans, Cheryl
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLECTIDES
TITLE OF INVENTION: ENCODING THEM
NUMBER OF SEQUENCES: 71
CORRESPONDENCE ADDRESS:
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SEQ ID NO 45
LENGTH: 265
TYPE: DNA
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APPLICANT: M1, Sha
APPLICANT: Genetics Institute, Inc.
APPLICANT: Genetics Institute, Inc.
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
FILE REFERENCE: 6006B.AJ172A
CURRENT APPLICATION NUMBER: US/09/175,928A
CURRENT APPLICATION NUMBER: US/09/175,928A
CURRENT FILING DATE: 1998-10-20
NUMBER OF SEQ ID NOS: 62
CORMANDE: Datantin Var. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Jacobs, Kenneth
APPLICANT: McCoy, John M.
APPLICANT: LaVallie, Edward R.
APPLICANT: Collins Racie, Lisa A.
APPLICANT: Evans, Cheryl
APPLICANT: Merberg, David
                                                NAME/KEY:
LOCATION:
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 46:
           NAME/KEY: unsure LOCATION: (133)
                                                                                                          FEATURE
                                                                                                                                                                                                                                                                                   NAME/KEY: unsure LOCATION: (79)
                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: unsure LOCATION: (2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Mus musculus FEATURE:
FEATURE:
                                                                                                                       LOCATION:
                                                                                                                                            NAME/KEY:
                                                                                                                                                                FEATURE
                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                    NAME/KEY: unsure
                                                                                                                                                                                                                                                                                                                                         NAME/KEY: unsure LOCATION: (75)
                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: unsure LOCATION: (67)
                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                             OCATION:
                                                                                                                                                                                                NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
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                                                                 unsure
(121)
                                                                                                                     unsure
(111)
                                                                                                                                                                           unsure
(104)
                                                                                                                                                                                                                                 (101)
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NAME/KEY: unsure LOCATION: (175)

EATURE

OCATION: (183)

NAME/KEY: unsure LOCATION: (172)

NAME/KEY: unsure LOCATION: (199)..(200)

NAME/KEY:

unsure (208)

VAME/KEY:

VAME/KEY: OCATION:

unsure (211)

NAME/KEY: unsure LOCATION: (192)

EATURE OCATION: EATURE

(187) unsure

PEATURE

FEATURE

OCATION: WAME/KEY: unsure

(164)

NAME/KEY: unsure LOCATION: (162)

FEATURE:

NAME/KEY: unsure LOCATION: (157)

FEATURE:

unsure (136)

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US-09-032-684-13/c
; Sequence 13, Application US/09032684
; Patent No. 5882874
                APPLICANT: Fisher, Paul B.
TITLE OF INVENTION: RECIPROCAL SUBTRACTION DIFFERENTIAL
TITLE OF INVENTION: DISPLAY
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper & Dunham LLP
1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                    Score 20; DB 4;
; Pred. NO. 4.9;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                      0; Indels
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US-08-967-101-82/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 82,
; Patent No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (212) 391-052 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 5555
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 278-0400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
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APPLICATION NUMBER: US/09/032,684
                                                                                                                                                                                                                      COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED
TITLE OF INVENTION: TO ALZHEIMER'S DISEASE
NUMBER OF SEQUENCES: 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 296 base pairs
                                                                                                                                         APPLICATION NUMBER: US/08/967,101 FILING DATE: 10-NOV-1997 CLASSIFICATION: 435 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
        NAME: Pitcher, Edmund R. TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 248-7000 TELEPAX: (617) 248-7100
                                                                                 FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 606 GNAAAAAAAAAAAAAAAAA 625
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: New York
STATE: New York
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CITY: F
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS
                                                                                                                                                                                                                                                                                                                                        STATE: Massachusetts
COUNTRY: U.S.A.
ZIP: 02110
                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2, Application US/08967101
5840540
                                                                                                                                                                                                                                                                                                                                                                                                   Boston
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High Street Tower - 125 High Street
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PatentIn Release #1.0, Version #1.30
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; NAME/KEY: unsure ; LOCATION: (246) US-09-175-928-45

FEATURE:

NAME/KEY: unsure LOCATION: (243)..(244)

OCATION:

unsure (227)

unsure (225) unsure (223)

Query Match 3.2%; Sometimes Similarity 100.0%; 1 Matches 20; Conservative 0;

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RESULT 9

GENERAL INFORMATION:

STREET:

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US-09-124-698-82/c

; Sequence 82, Application US/09124698

; Patent No. 6117978
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US-08-592-541-82/c
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                                                                                                                                                                                                                                                                                            TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 82
SEQUENCE CHARACTERISTICS:
LENGTH: 500 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                              Query Match
                                                                                                                                                                                     Matches
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 82, Application US/08592541 Patent No. 5986054
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                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: Pitcher, Edmund R.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: TESTA, HURWITZ & THIBEAULT
STREET: High Street Tower - 125 High Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: GENETIC SEQUENCES AND ITITLE OF INVENTION: TO ALZHEIMER'S DISEASE NUMBER OF SEQUENCES: 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
                                                                                                                                                                                                                                                                MOLECULE TYPE: DNA (genomic)
                                                                                                                                              606 GNAAAAAAAAAAAAAAAAA 625
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                                                                                                                                                                                                Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
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                                                                                                                                                                                                Similarity
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                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                          Sequence 82, Application US/09127480 Patent No. 6194153 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 82:
           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS: TESTA, HU
                                                                                                                                                                                                                                                                       APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: ERASER, PAUL E
TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED
TITLE OF INVENTION: TO ALZHEIMER'S DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Pitcher, Edmund R. TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, JOHANNA M
APPLICANT: FRASER, PAUL E
TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED
NUMBER OF SEQUENCES: 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                606 GNAAAAAAAAAAAAAAAA 625
                                                                                                                                                                              STREET: High
CITY: Boston
                                                                                                                                  ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          34 GNAAAAAAAAAAAAAAAA 15
APPLICATION NUMBER:
                                                                                                                                                                    STATE: Massachusetts
                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 500 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20;
                                                                                                                                  02110
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                                                                                                                                                                                                     E: TESTA, HURWITZ & THIBEAULT
High Street Tower - 125 High
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High Street Tower - 125 High Street
                                                                                                                                                  U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Floppy disk
                                                                                                                                                                                                                                                                 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.2%; Score 20; DB 3; Length 500; 100.0%; Pred. No. 4.7;
US/09/127,480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US/09/124,698
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                                                                                                                                                                                                   125 High Street
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CLASSIFICATION:
PRIOR APPLICATION DATA:

APPLICATION NUMBER:

US/08/592,541

FILING DATE:

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US-08-496-841C-82/c
Sequence 82, Application US/08496841C
Patent No. 6210919
GENERAL INFORMATION:
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                                                  SEQUENCE DESCRIPTION: SEQ ID NO: 82: US-08-496-841C-82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
 Best Local Similarity
                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (617) 248-7100 INFORMATION FOR SEQ ID NO: 82:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Pitcher, Edmund R.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 82: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS
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STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/496,841C
FILING DATE: 28-Jun-1995
CLASSIFICATION: <UNKnown>
ATTORNEY/AGENT INFORMATION:
NAME: Paul F. Febliner, Ph.D.
REGISTRATION NUMBER: 35,135
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Darby & Darby, PC
                                                                                        TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: ST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20;
                                                                                                                         LENGTH: 500 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 10022
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 805 Third Avenue
                                                                                                                                                                                                                    TELEPHONE: (212) 527-7700
TELEFAX: (212) 753-6237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            500 base pairs
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FRASER, PAUL E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     U.S.A.
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100.0%; Prr
3.2%;
100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          JOHANNA M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   175
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Pred. No.
Score 20; DB 4; Length 500; Pred. No. 4.7;
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US-09-124-523-82/c
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; Patent No. 6
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US-09-020-956-32/c
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                                                                                                                                                              Sequence 32, Application US/09020956 Patent No. 6261562
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                                                                   GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillin, Davin C.
APPLICANT: Dillin, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND METHODS
NUMBER OF SEQUENCES: 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED
TITLE OF INVENTION: TO ALZHEIMER'S DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Pitcher, Edmund R. TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 248-7000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
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LENGTH: 500 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                  CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
STREET: 6300
CITY: Seattle
               ADDRESSEE: SEED and BERRY LLP STREET: 6300 Columbia Center,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20; Conservative
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                                                                                                                                                                                                                                                                                                                                                    Conservative
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                   701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                         DB 4; Length 500;
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us-09-030-607-32/c
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                                                       NAME: MAKI, DAVIG J

REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 2101:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 32:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 32, Application US/09030607 Patent No. 6262245
GENERAL INFORMATION:
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LENGTH: 789 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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REFERENCE/DOCKET NUMBER: 210
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 32:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09
FILING DATE: 25-FEB-1998
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Xu, Jiangchun APPLICANT: Dillon, Davin C.
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CURRENT APPLICATION DATA:
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NAME: Maki, David J.
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                                 LENGTH:
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STREET: 6300 Columbia Center, 701 Fifth Avenue
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COUNTRY:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Local Similarity 100.0%; P
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nucleic acid
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GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun

APPLICANT: Dillon, Davin C.

APPLICANT: Mitcham, Jennifer L.

APPLICANT: Harlocker, Susan Louise

APPLICANT: Jiang Yuqui

APPLICANT: Reed, Steven G.
                                                                                                                                                           US-09-439-313-32/c
                                                                                                                                                                                                                                                                                                                                                               ; NAME/KEY: misc_feature COCATION: (1)...(789) ; OTHER INFORMATION: n = A,T,C or G US-09-605-785-32
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US-09-605-785-32/c
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                                                                                                                      Sequence 32, Application US/09439313 Patent No. 6329505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Wang, Aijun
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 20121.427C16
CURRENT APPLICATION NUMBER: US/09/605,785
CURRENT FILING DATE: 2000-06-27
NUMBER OF SEQ ID NOS: 835
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
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Best Local S
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Best Local Similarity
Matches 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Homo sapien
FEATURE:
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TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                        ch 3.2%; Score 20; DB 4; Length 789; I Similarity 100.0%; Pred. No. 4.5; 20; Conservative 0; Mismatches 0; Indels
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Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
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ick, Thomas S.
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RESULT 20
US-09-352-616A-32/c
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; LCCATION: (1)...(789)
; OTHER INFORMATION: n = A,T,C or G
US-09-439-313-32
                          RESULT 21
US-09-232-149A-32/c
; Sequence 32, Application US/09232149A
; Patent No. 6465611
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: COMPOSITIONS OF PROSTATE CANCER
TITLE OF INVENTION: DAY, OF PROSTATE CANCER
TITLE OF INVENTION: COMPOSITION OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
CURRENT FILING DATE: 1999-11-12
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
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APPLICANT: Dillon, Davin C.
APPLICANT: Harlocker, Susan Louise
APPLICANT: Jiang, Yuqui
APPLICANT: Jiang, Yuqui
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jennifer Lynn
APPLICANT: INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF PROSTATE CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.427C8
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                GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                  Query Match
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CURRENT FILING DATE: 1999-07-13
NUMBER OF SEO ID NOS: 472
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APPLICANT: Xu, Jiangchun
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ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 789
TYPE: DNA
ORGANISM: Homo sapien
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LOCATION: (1)...(789)
OTHER INFORMATION: n =
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Fanger, Gary
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100.0%; Pred. No. 4.9
ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 1998-09-08
EARLIER APPLICATION NUMBER: PCT/US98/04493
EARLIER FILING DATE: 1998-03-06
EARLIER APPLICATION NUMBER: 60/040,162
EARLIER FILING DATE: 1997-03-07
EARLIER APPLICATION NUMBER: 60/040,333
EARLIER FILING DATE: 1997-03-07
EARLIER APPLICATION NUMBER: 60/038,621
EARLIER APPLICATION NUMBER: 60/038,621
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Dillon, Davin C.
APPLICANT: Mitcham, Jennifer Lynn
APPLICANT: Mitcham, Jennifer Lynn
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE
TITLE OF INVENTION: CANCER AND METHODS FOR THEIR USE
TILE REFERENCE: 210121.427C6
CURRENT APPLICATION NUMBER: US/09/232,149A
CURRENT FILING DATE: 1999-01-15
NUMBER OF SEQ ID NOS: 338
SOFTWARE: PASTSEQ for Windows Version 3.0
SEQ ID NO 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL
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                                                                                                                                                                                                                                                EARLIER APPLICATION NUMBER: 60/047,615
EARLIER FILING DATE: 1997-05-23
EARLIER APPLICATION NUMBER: 60/047,597
EARLIER FILING DATE: 1997-05-23
EARLIER APPLICATION NUMBER: 60/047,502
EARLIER FILING DATE: 1997-05-23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: 186 Human Secreted proteins
                                                            EARLIER APPLICATION NUMBER: 60/047,617
EARLIER FILING DATE: 1997-05-23
EARLIER APPLICATION NUMBER: 60/047,618
EARLIER FILING DATE: 1997-05-23
                                                                                                                                                                                                                                                                                                                                                                                        EARLIER APPLICATION NUMBER: 60/040,163
EARLIER FILING DATE: 1997-03-07
EARLIER APPLICATION NUMBER: 60/047,600
EARLIER FILING DATE: 1997-05-23
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EARLIER FILING DATE: 1997-03-07
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LOCATION: (1)...(789)
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA ORGANISM: Homo sapien
                                                                                                                                                                                                    EARLIER
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EARLIER FILING DATE: 1997-03-07
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                                                                                                                                                      FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,583
FILING DATE: 1997-05-23
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                                                                                                                                                                                                                             APPLICATION NUMBER: 60/047,633
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                 APPLICATION NUMBER: 60/047,503 FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,592
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.00.0%; Pred. No. 4.5
.ve 0; Mismatches
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2R FILING DATE: 1997-08-22
2R APPLICATION NUMBER: 60/056,872
2R FILING DATE: 1997-08-22
2R PELICATION NUMBER: 60/056,882
2R APPLICATION NUMBER: 60/056,682
2R APPLICATION NUMBER: 60/056,637
2R FILING DATE: 1997-08-22
2R APPLICATION NUMBER: 60/056,903
2R FILING DATE: 1997-08-22
2R APPLICATION NUMBER: 60/056,903
2R FILING DATE: 1997-08-22 R FILING DATE: 1997-04-11

R APPLICATION NUMBER: 60/043,315

R FILING DATE: 1997-04-11

R APPLICATION NUMBER: 60/048,974

R FILING DATE: 1997-06-06

R APPLICATION NUMBER: 60/056,886

R APPLICATION NUMBER: 60/056,886 JR APPLICATION NUMBER: 60/043,669

IR FILLING DATE: 1997-04-11

PR APPLICATION NUMBER: 60/043,312

PR FILLING DATE: 1997-04-11

JR APPLICATION NUMBER: 60/043,313

REFILLING DATE: 1997-04-11 IR APPLICATION NUMBER: 60/047,632
IR FILLING DATE: 1997-05-23
IR APPLICATION NUMBER: 60/047,601
IR FILLING DATE: 1997-05-23
IR FILLING DATE: 1997-04-11
IR FILLING DATE: 1997-04-11 FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,893
FILING DATE: 1997-08-22 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,878 APPLICATION NUMBER: 60/056,630 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/U APPLICATION NUMBER: 60/056,877 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/043,672 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,671 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,674 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,314
FILING DATE: 1997-04-11
APPLICATION NUMBER: 60/043,569
FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,311 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,568 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/047,598
FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,613 APPLICATION NUMBER: 60/047,612 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,596 FILING DATE: 1997-05-23 FILING DATE: 1997-0 APPLICATION NUMBER: APPLICATION NUMBER: 60/ FILING DATE: 1997-05-23 APPLICATION NUMBER: FILING DATE: 1997-0: APPLICATION NUMBER: 6 FILING DATE: 1997-05-APPLICATION NUMBER: 60 FILING DATE: 1997-05-FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/ FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 1997-08-22 1997-05-23 1997-05-23 60/056,889 60/047,582 60/047,587 60/047,500 60/047,492 60/047,584 60/047,581

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R APPLICATION NUMBER: 60/01
R FILING DATE: 1997-08-22
R APPLICATION NUMBER: 60/01
R FILING DATE: 1997-08-22 R FILING DATE: 1997-05-23

R APPLICATION NUMBER: 60/043,670

R FILING DATE: 1997-04-11

R APPLICATION NUMBER: 60/056,632

R FILING DATE: 1997-08-22

R APPLICATION NUMBER: 60/056,664

R FILING DATE: 1997-08-22 R APPLICATION NUMBER: 60/047,614
IR FILING DATE: 1997-05-23
IR APPLICATION NUMBER: 60/043,578
IR FILING DATE: 1997-04-11
IR APPLICATION NUMBER: 60/043,576
IR FILING DATE: 1997-04-11 PR APPLICATION NUMBER: 60/047,586

RE FILLING DATE: 1997-05-23

PR APPLICATION NUMBER: 60/047,590

PR FILLING DATE: 1997-05-23

PR APPLICATION NUMBER: 60/047,594

PR FILLING DATE: 1997-05-23

PR APPLICATION NUMBER: 60/047,589

PR FILLING DATE: 1997-05-23

PR APPLICATION NUMBER: 60/047,589

PR FILLING DATE: 1997-05-23

PR APPLICATION NUMBER: 60/047,593

PR APPLICATION NUMBER: 60/047,593 APPLICATION NUMBER: 60/056,887 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,908 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/048,964 APPLICATION NUMBER: 60/ APPLICATION NUMBER: 60/056,876 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,881 FILING DATE: FILING DATE: 1997-05-23
APPLICATION NUMBER: 60/047,588
FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,501 APPLICATION NUMBER: 60/ FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/ APPLICATE: 1997-08-22
APPLICATION NUMBER: 60/056,864
APPLICATION 1997-08-22 APPLICATION NUMBER: 60/047,585 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/ APPLICATION NUMBER: 60/057,761 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,892 FILING DATE: 1997-08-22 FILING DATE: 1997-0
APPLICATION NUMBER: APPLICATION NUMBER: 60/0 APPLICATION NUMBER: 60/056,874 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/ FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,888 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,879 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,636 APPLICATION NUMBER: FILING DATE: 1997-08-FILING DATE: APPLICATION NUMBER: 60/056,880 FILING DATE: 1997-08-22 1997-08-22 1997-08-22 60/056,862 60/056,875 60/056,909 60/047,599 60/047,595 60/056,845 60/056,631 60/056,911 60/056,894

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US-09-443-041A-29
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                                  GENERAL INFORMATION:
APPLICANT: Robert G. Kuimelis
APPLICANT: Robert G. Kuimelis
TITLE OF INVENTION: METHODS FOR CODING AND SORTING IN VITRO
TITLE OF INVENTION: TRANSLATED PROTEINS
FILE REFERENCE: 50036/032002
CURRENT APPLICATION NUMBER: US/09/648,040
CURRENT FILING DATE: 2000-08-25
PRIOR APPLICATION NUMBER: US 60/151,261
PRIOR FILING DATE: 1999-08-27
NUMBER OF SEQ ID NOS: 11
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Best Local (
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LENGTH: 1210
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EARLIER FILING DATE: 1997-08-22
EARLIER APPLICATION NUMBER: 60/057,669
EARLIER FILING DATE: 1997-09-05
EARLIER APPLICATION NUMBER: 60/049,610
EARLIER FILING DATE: 1997-06-13
EARLIER FILING DATE: 1997-06-13
EARLIER APPLICATION NUMBER: 60/061,060
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SEQ ID NO 4
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CURRENT FILING DATE: 1999-11-18
FRIOR APPLICATION NUMBER: 60/109,283
PRIOR ETLING DATE: 1998-11-20
NUMBER OF SEQ ID NOS: 33
SOFTWARE: Microsoft Office 97
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EARLIER APPLICATION NUMBER: 60/057,650
EARLIER FILING DATE: 1997-09-05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EARLIER FILING DATE: 1997-10-02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Sterol Metabolism Enzymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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                      SOFTWARE: FastSEQ for Windows Version 4.0
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LOCATION: 1129
OTHER INFORMATION: any nucleotide
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ORGANISM: Triticum aestivum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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Rafalski, Antoni
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; NAME/KEY: misc_feature
; LOCATION: 40
; OTHER INFORMATION: /product= "N is A,
US-08-859-106A-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 4, Application US/08859106A Patent No. 5965422
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LOCATION: 10
OTHER INFORMATION: n at position 10 can be a, t, c, or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 30
                                                                                                                                                                                               TELEFAX: (703) 836-2021 INFORMATION FOR SEQ ID NO: 4: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6620
                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: DE 19
FILING DATE: 22-MAY-1996
ATTORNEY_AGENT INFORMATION:
NAME: McGowan, Malcolm K.
REGISTRATION NUMBER: 39,3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: LYSOPHOSPHOLIPASE PRODUCED FROM TITLE OF INVENTION: ASPERGILLUS BY RECOMBINANT METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                    FEATURE:
                                                                                  ANTI-SENSE:
                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               607 ИАААААААААААААААА 625
                                                                                                 HYPOTHETICAL:
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                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
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                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                LENGTH:
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                                                                                                                                                               nucleotide
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                                                                                                                                                                                40 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P.O. Box 1404
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Nerven, Quoc Khanh
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                                  misc_feature
                                                                                                                                                                                                                                (703) 836-2021
                                                                                   Z
                                                                                                                                  linear
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                                                                                                 genomic DNA
No
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20-MAY-1997
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16;
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CURRENT APPLICATION NUMBER: US/09/011,540
CURRENT FILING DATE: 1998-04-20
EARLIER APPLICATION NUMBER: DE 19526485.1
EARLIER FILING DATE: 1995-07-20
EARLIER APPLICATION NUMBER: PCT/EP96/01430
EARLIER APPLICATION NUMBER: PCT/EP96/01430
EARLIER FILING DATE: 1996-04-01
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 11
EARLIER FILING DATE: 1996-04-01
SEQ ID NO 11
EARLIER FILING DATE: 12
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                 SEQ ID NO 5
LENGTH: 51
                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                    APPLICANT: WEINDEL, Kurt
APPLICANT: SEIDEL, Cristoph
APPLICANT: LASSONCZYK, Gerhard
TITLE OF INVENTION: METHOD OF DETECTING A SUBSTANCE TO BE ANALYZED
FILE REFERENCE: 101614-07096
CURRENT APPLICATION NUMBER: US/09/051,079A
CURRENT FILING DATE: 1998-09-10
EARLIER APPLICATION NUMBER: PCT/EEP96/04358
EARLIER APPLICATION NUMBER: DE/195 37 952.7
EARLIER FILING DATE: 1996-10-12
EARLIER FILING DATE: 1995-10-12
                                                                                                                                                                                                                                                                                                                                           Sequence 5, Application US/09051079A Patent No. 6214549
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                                                   SOFTWARE: PatentIn Ver. 2.1
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APPLICANT: WALF, Sabine
APPLICANT: PLAINER, HERMANN
TITLE OF INVENTION: LEUCINE AMINOPEPTIDASES PRODUCED RECOMBINANTLY FROM
TITLE OF INVENTION: ASPERGILLUS SOYAE
FILE REFERENCE: 015200-051
TYPE: DNA
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NAME/KEY: misc_feature
LOCATION: (39)...(40)
OTHER INFORMATION: N is A, C or G.
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APPLICANT: GOTTSCHALK, Michael
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          607 NAAAAAAAAAAAAAAAAA 625
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100.0%; Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 4; Length 40; 16;
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RESULT 28
US-08-618-100B-9/c
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INFORMATION FOR SEQ ID NO:
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                                              REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 211
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                      OVERANTING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/618,100B
FILING DATE: March 19, 1996
CLASSIFTANTON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                            FILING DATE: October 30, 1995
APPLICATION NUMBER: 08/510,584
FILING DATE: August 2, 1995
APPLICATION NUMBER: 08/418,096
FILING DATE: APPLI 5, 1995
APPLICATION NUMBER: 08/408,584
FILING DATE: March 20, 1995
                                                                                                                                                                                                                                                                            CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Oligodesoxyribonucleotide
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APPLICANT: Miller, Stephen G.
APPLICANT: MILLER, Stephen G.
APPLICANT: MODULATORS OF Ob GENE AND
APPLICANT: MODULATORS OF THEREFO.
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NAME/KEY: misc_feature
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          TELEPHONE: (213) 489-160
TELEFAX: (213) 955-0440
TELEX: 67-3510
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                607 NAAAAAAAAAAAAAAAAA 625
                                                                                             REGISTRATION NUMBER:
                                                                                                                                                                                                           FILING DATE: October APPLICATION NUMBER: FILING DATE: Augus
                                                                                                                                                                                                                                              APPLICATION NUMBER: 0 FILING DATE: October
                                                                                                                    NAME:
                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: 3.5" Di
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5, 6068976
                                                                                                             Warburg, Richard J.
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California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Suite 4700
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633 West Fifth Street
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Staels, Bart
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                              IBM Compatible
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5, 1995
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US-08-702-344-7
В
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                                                                                                                         US-08-702-344-7
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                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
TELEPHONE: (617) 876-5851
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 69 base pairs
                                                                                           Query Match
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LENGTH: 52 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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APPLICANT:
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APPLICANT: Merberg, Maurice
APPLICANT: Treacy, Maurice
APPLICANT: Spaulding, Vikki
APPLICANT: SPAULDING FOR AND POLYNUCLEOTIDES
APPLICANT: SECRETED PROTEINS AND POLYNUCLEOTIDES
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CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  607 NAAAAAAAAAAAAAAAA 625
                                                                                                                                        MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    51 NAAAAAAAAAAAAAAAA 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY:
STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: "N" represents any base
                            607 ИАЛАЛАЛАЛАЛАЛАЛАЛАЛ 625
                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 536
                                                          Local Similarity
les 19; Conserv
 5 NAAAAAAAAAAAAAAA 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/08702344
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LaVallie, Edward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         U.S.A.
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                                                           3.0%; Scilarity 100.0%; For Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Genetics Institute, Inc.
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Pred. No.
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                                                                  Mismatches
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15;
                                                                               DB 1; Length 69; 15;
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RESULT 30
US-08-702-344-22
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US-09-269-911A-7/c
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                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                     sequence 7, Application US/09269911A Patent No. 6228589
                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                           SOFTWARE:
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                                                                                                                                                                                                                                                                                                                                                                                                   Best
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INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
                                                                                                                                         APPLICANT: Brenner, Sydney
TITLE OF INVENTION: MEASUREMENT OF GENE EXPRESSION PROFILES
TITLE OF INVENTION: IN TOXICITY-DETERMINATION
FILE REFERENCE: 5525-0025.10
                                                                                      CURRENT APPLICATION NUMBER: US/09/269,911A
CURRENT FILING DATE: 2000-02-28
NUMBER OF SEO ID NOS: 10
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APPLICANT: Treacy, Maurice
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES
TITLE OF INVENTION: ENCODING THEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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TELECOMMUNICATION INFORMATION:
           TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: cDNA
                                            ENGTH: 69
                                                                                                                                                                                                                                                                                                                                                607 NAAAAAAAAAAAAAAAAA 625
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TOPOLOGY: li
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Massachusetts
                                                                             FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
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100.0%; Pr
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Pred. No.
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                                                                                             Query Match
                                                                Matches
                                                                                Best
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                                                                                                                                                                                                                                    TELEFAX: (617) 876-5851 INFORMATION FOR SEQ ID NO:
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APPLICANT:
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NAME/KEY: misc_feature
LOCATION: (65)...(65)
OTHER INFORMATION: n = cDNA from library
                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/664,596B
                                                                                                                                                                                                                                                                  NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES
TITLE OF INVENTION: ENCODING THEM
                             607 NAAAAAAAAAAAAAAAAA 625
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                                                                             Local
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CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
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8 NAAAAAAAAAAAAAAAA 26
                                                                                                                                                                                                     LENGTH:
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                                                                           Similarity
                                                                                                                                                                                    nucleic acid
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                                                                                                                                                                                                   84 base pairs
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87 CambridgePark Drive
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Treacy, Maurice
Evans, Cheryl
Spaulding, Vikki
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LaVallie, Edward
                                                              Conservative
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                                                                                                                                                                      double
                                                                           3.0%; Score 19;
100.0%; Pred. No.
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Pred. No.
                                                           Mismatches
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; TOPOLOGY: li
; MOLECULE TYPE:
US-08-738-367-3
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US-08-991-789A-266/c
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US-08-738-367-3
                                                                                        Sequence 266, Application US/08991789A
Patent No. 6225054
GENERAL INFORMATION:
Smith, John M.
Smith, John M.
Reed, Steven G.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TREATMENT AND DIAGNOSIS OF BREAST CANCER
                                                                                                                                                                                                                                                                                                                                                                          Matches
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Patent No. 5827688
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INFORMATION FOR SEQ ID NO: 3:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/738,367
FILING DATE:
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APPLICANT:
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APPLICANT:
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LENGTH: 84 base pairs
TYPE: nucleic acid
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
APPLICANT: Spaulding, vikki
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES
TITLE OF INVENTION: ENCODING THEM
                                                                                                                                                                                                                                                                                                                                    607 НААААААААААААААААА 625
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STATE: Massachu
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                                             NUMBER OF SEQUENCES: 292
CORRESPONDENCE ADDRESS:
ADDRESSEE: Seed IP Law Group
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02140
                 CITY: Seattle
                              STREET: 701 Fifth Avenue, Suite 6300
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LaVallie, E
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Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           double
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100.0%; Pred. No.
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COUNTRY: USA

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US-09-062-451-266/c
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                                          TELEFAX: (206) 682-60 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 266:
SEQUENCE CHARACTERISTICS:
LENGTH: 193 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
REFERENCE/DOCKET NUMBER: 210
FELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION UMBER: US/09/062,451 FILING DATE: 04-APR-1997
                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 297
CORRESPONDENCE ADDRESS:
ADDRESSE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Frudakis, Tony N. APPLICANT: Smith, John M. APPLICANT: Reed, Steven G.
                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                          CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TITLE OF INVENTION: TREATMENT AND DIAGNOSIS OF BREAST CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     180 NAAAAAAAAAAAAAAAAAA 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           607 NAAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Seattle
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PStephin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/991,789A
FILING DATE: 11-Dec-1997
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: POTTER, Jane E. R.
REGISTRATION NUMBER: 33,332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19;
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193 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                               266:
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                                                           Query Match
Best Local Similarity
Matches 19; Conserv
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                                                                                                                                                                                                                                       TELEFAX: (617) 876-58 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, Inc
STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
APPLICANT: Evans, Cheryl
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED PROTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                         TOPOLOGY: 1: MOLECULE TYPE:
                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 208 base pairs
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                            607 NAAAAAAAAAAAAAAAAA 625
161 NAAAAAAAAAAAAAAAAA 179
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STRANDEDNESS: sing
TOPOLOGY: linear
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les 19; Conserv
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                                                                                                                                                                        STRANDEDNESS:
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(617) 876-5851
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LaVallie, Edward
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                                                             Conservative
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                                                                                                                                                           linear
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/ 100.0%; Pr
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                                                                            100.0%;
                                                                            3.0%; Score 19; DB 1; 100.0%; Pred. No. 14;
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                                                             0;
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                                                              Mismatches
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                                                              0; Indels
                                                                                              Length 208;
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RESULT 37 US-08-967-101-98/c ; Sequence 98, Application US/08967101

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; TOPOLOGY: linear ; MOLECULE TYPE: DNA (genomic) US-08-967-101-98
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US-08-592-541-98/c
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Best Local Similarity
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (617) 248-700
TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Pitcher, Edmund R.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED TITLE OF INVENTION: TO ALZHEIMER'S DISEASE NUMBER OF SEQUENCES: 183
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 208 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                 APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
                                                                                                                                                                                                                                           TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED TITLE OF INVENTION: TO ALZHEIMER'S DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            159 NAAAAAAAAAAAAAAAA 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             607 NAAAAAAAAAAAAAAAAA 625
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CITY: Boston
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/01 FILING DATE: 10-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                            COUNTRY:
                                                                                                                                                   CITY: Boston
                                                                                                                                                                   STREET:
                                                                                                                                                                                    ADDRESSEE:
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                                                                                                                                 Massachusetts
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                                                                                                                                                                   E: TESTA, HURWITZ & THIBEAULT
High Street Tower - 125 High Street
                                                                                                              U.S.A.
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PatentIn Release #1.0, Version #1.30
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100.0%; Pred. No.
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14;
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US-09-124-698-98/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: ST. GEORGE
APPLICANT: ROMMENS, J
APPLICANT: FRASER, PA
TITLE OF INVENTION: G
TITLE OF INVENTION: G
TITLE OF INVENTION: G
TITLE OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 98, Application US/09124698 Patent No. 6117978
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                 Query Match
 Best Local Similarity
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LEGGTH: 208 base pairs
                                                                                                                                                                   TELEFAX: (617) 248-7100 INFORMATION FOR SEQ ID NO: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/592,541
                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Pitcher, Edmund R.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEB: TESTA, HURWITZ & THIBEAULT
STREET: High Street Tower - 125 High Street
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                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 208 base pairs
                                                                 MOLECULE TYPE: DNA (genomic)
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STRANDEDNESS: single
TOPOLOGY: linear
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Boston
STATE: Massachusetts
                                                                                      TOPOLOGY:
                                                                                                 STRANDEDNESS:
                                                                                                                     TYPE:
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                                                                                     linear
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                                                                                                     single
3.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     183
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 Score 19;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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 DB
14;
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                   Length 208;
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RESULT 41
US-08-496-841C-98/C
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US-09-127-480-98/c
                                                                                                                                         Sequence 98, Application US/08496841C Patent No. 6210919
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                             Matches
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (617) 248-7100 INFORMATION FOR SEQ ID NO: 98:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Pitcher, Edmund R. TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 248 7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 208 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED
TITLE OF INVENTION: TO ALZHEIMER'S DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
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ADDRESSEE: TESTA, HURWITZ &
STREET: High Street Tower -
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                       159 NAAAAAAAAAAAAAAAAA 141
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                        3.0%; Score 19; Local Similarity 100.0%; Pred. No. es 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
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                                                                                                       APPLICANT: ST. GEORGE-HYSLOP, PETER H
                  CORRESPONDENCE ADDRESS:
                                      NUMBER OF SEQUENCES:
                                                   TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED TO ALZHEIMER'S DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98, Application US/09127480
o. 6194153
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ADDRESSEE: Darby & Darby, PC
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                                                                                          FRASER, PAUL E
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125 High Street
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US-09-175-928-37, Application US/09175928A
: Patent No. 6312921
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; MOLECULE TYPE: DNA (genomic);
SEQUENCE DESCRIPTION: SEQ ID NO: 98:
US-08-496-841C-98
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                                                                                                                                                                                             SEQ ID NO 37
LENGTH: 208
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                     APPLICANT: Evans, Cheryl
APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
APPLICANT: Mi, Sha
APPLICANT: Mi, Sha
APPLICANT: Genetics Institute, Inc.
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
FILE REFERENCE: 6006B.AJ172A
                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Jacobs, Kenneth APPLICANT: McCoy, John M. APPLICANT: LAVAILIE, Edward R.
                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/175,928A CURRENT FILING DATE: 1998-10-20
                                                                                                                                                                                                                                                  SOFTWARE:
                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
              NAME/KEY:
                                                                                                                                                           ORGANISM: Mus musculus FEATURE:
                                                   FEATURE:
                                                                                     FEATURE:
NAME/KEY: unsure
                                                                                                                      NAME/KEY: unsure LOCATION: (8)
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EATURE:
                                                                    CATION:
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APPLICATION NUMBER: US/08/496,841C
FILING DATE: 28-Jun-1995
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Paul F. Fehlner, Ph.D.
REGISTRATION NUMBER: 35,135
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 527-7700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                 208
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                                                                                                                                                                                                                                                  PatentIn Ver. 2.0
              unsure
(31)
                                                                  (29)
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14;
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US-09-124-523-98/c

Sequence 98, Application US/09124523

Patent No. 6395960

GENERAL INFORMATION:

APPLICANT: ST. GEORGE-HYSLOP, PETER H

APPLICANT: FRASER, PAUL E

TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEIN

TITLE OF INVENTION: TO ALZHEIMER'S DISEASE

NUMBER OF SEQUENCES: 183

CORRESPONDENCE ADDRESS:

ADDRESSEE: TESTA, HURWITZ & THIBEAULT

STREET: High Street Tower - 125 High Street
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                                                                                                                                                                                                    RESULT 43
                                                                                                                                                                                                                                                                                                     Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: unsure
LOCATION: (161)
                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: unsure LOCATION: (137)
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LOCATION: (133)
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LOCATION: (130)..(131)
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LOCATION: (100)..(101)
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LOCATION: (91)
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OCATION: (86)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AME/KEY: unsure
                                                                                                                                                                                                                                              161 NAAAAAAAAAAAAAAAAA 179
                                                                                                                                                                                                                                                                       607 NAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                                                                                                                                    Local Similarity
les 19; Conserv
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(128)
Boston
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100.0%; Pr
                                                       GENETIC SEQUENCES AND PROTEINS RELATED TO ALZHEIMER'S DISEASE
                                                                                                                                                                                                                                                                                                                 Score 19; DB 4; Length 208; Pred. No. 14;
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Best Local S
Matches 19
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Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Pitcher, Edmund R.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
TELEFAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 98:
SEQUERICE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                APPLICANT: Merberg, David
APPLICANT: Treacy, Maurice
APPLICANT: Evans, Cheryl
APPLICANT: Spaulding, Vikki
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES
TITLE OF INVENTION: ENCODING THEM
NUMBER OF SEQUENCES: 71
CORRESPONDENCE ADDRESS:
               CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Brown, Scott A.
                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
                                                                                                                                                                                                      COUNTRY: U
ZIP: 02140
                                                                                                                                                                                                                                                                    ADDRESSEE:
STREET: 8
REGISTRATION NUMBER: 32,724
                                                                 APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                    STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH:
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19; Conserv
                                                                                                                                                                                                                                                      Cambridge
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                                                                                                                                                                                                                   Massachusetts
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87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    208 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                         McCoy, John
LaVallie, E
                                                                                                                                                                                                                                                                                                                                                                                                                                           Racie, Lisa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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100.0%; Pr
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; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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APPLICANT: Endege, Wilson O.
APPLICANT: Steinmann, Kathleen E.
APPLICANT: Astle, Jon H.
APPLICANT: Astle, Jon H.
APPLICANT: Burgess, Christopher C.
APPLICANT: Bushnell, Steven E.
APPLICANT: Bushnell, Steven E.
APPLICANT: Carroll III, Eddie
APPLICANT: Carroll III, Eddie
APPLICANT: Carroll III, Eddie
APPLICANT: Carroll III, Eddie
APPLICANT: Ford, Donna M.
APPLICANT: Monahan, John E.
APPLICANT: Monahan, John E.
APPLICANT: Schlegel, Robert
TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
TITLE OF INVENTION: MOVEL HUMAN
TITLE OF INVENTION: NOWBER: US/09/328,111
CURRENT APPLICATION NUMBER: US/09/328,111
CURRENT FILING DATE: 1999-06-08
EARLIER FILING DATE: 1999-06-08
EARLIER FILING DATE: 1998-06-10
NUMBER OF SEQ ID NOS: 850
SOFTMARE: FastSEQ for Windows Version 3.0
SEQ ID NO 715
LENGTH: 310
TYPE: DNA
ORGANISM: Homo Sapiens
FEATURE:
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US-09-328-111-715
US-09-328-111-715
; Sequence 715, Application US/09328111
; Patent No. 6262333
; GENERAL INFORMATION:
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NAME/KEY: misc_feature

LOCATION: (1)...(310)

OTHER INFORMATION: n = A,T,C or G
US-09-328-111-715
Search completed: November 7, Job time: 31.4202 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
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INFORMATION FOR SEQ ID NO: 43:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 244 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compugen Ltd

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Result
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Minimum DB seq length: 0
Maximum DB seq length: 200000000
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/cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
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8163.686 Million cell updates/sec
  Sequence 210, App Sequence 214, App Sequence 205, App Sequence 2011, App Sequence 2111, App Sequence 2409, App Sequence 2409, App Sequence 1516, App Sequence 1532, App Sequence 1532, App Sequence 1283, App Sequence 1283, App Sequence 698, App Sequence 698, App Sequence 379, App Sequence 379, App Sequence 379, App Sequence 379, App Sequence 539, App Sequence 539, App Sequence 539, App
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|   | 45                | 44                | 43                | 42                | 41                 | 40                | 39                | 38   | 37     | 36               | 35               | 34       | <b>3</b> 3       | 32               | 31                | 30                  | 29                | 28                | 27               | 26                | 25                | 24                | 23                | 22                | 21                 | 20                |
| ٠ | 19                | 19                | 19                | 19                | 19                 | 19                | 19                | 19   | 20     | 20               | 20               | 20       | 20               | 20               | 20                | 20                  | 20                | 20                | 20               | 20                | 20                | 20                | 20                | 20                | 20                 | 21                |
|   | 3.0               | 3.0               | 3.0               | 3.0               | 3.0                | 3.0               | 3.0               | 3.0  | 3.2 3  | 3.2              | 3.2              | 3.2      | 3.2              | 3.2              | 3.2               | 3.2                 | 3.2               | 3.2               | 3.2              | 3.2               | 3.2               | 3.2               | 3.2               | 3.2               | 3.2                | 3.4               |
|   | 60                | 56                | 56                | 56                | 53                 | <u>წ</u>          | 52                | 51   | 368004 | 789              | 789              | 789      | 789              | 789              | 648               | 496                 | 451               | 291               | 265              | 226               | 217               | 97                | 66                | 60                | 58                 | 1402              |
|   | 10                | 10                | 10                | 10                | 10                 | 10                | 10                | 10   | 10     | 10               | 10               | 10       | 10               | 10               | 10                | 10                  | 10                | 10                | 12               |                   | 10                | 10                | 10                | 10                | 10                 | 10                |
|   | US-09-925-301-758 | US-09-919-580-713 | US-09-919-580-677 | US-09-919-580-171 | US-09-783-590-3220 | US-09-919-580-705 | 783               | .783 | 49-654 | US-09-115-453-32 | US-09-822-827-32 | )30-606- | US-09-780-669-32 | US-09-759-143-32 | US-09-925-301-819 | US-09-783-590-10830 | US-09-777-564-625 | US-09-834-975-374 | US-10-040-916-46 | US-10-046-935-291 | US-09-895-828-248 | US-09-815-343-241 | US-09-925-301-759 | US-09-919-580-419 | US-09-783-590-2406 | US-09-925-297-171 |
|   | -                 | Sequence 713, App |                   | •                 | Ö                  | Sequence 705, App | Sequence 5881, Ap | 808  | ω<br>Α | 32,              | 32,              | 32,      |                  | 32,              | 819,              | 3                   | 625,              | 374,              |                  | 291, 🌶            | Sequence 248, App |                   | 759,              | Sequence 419, App | ō                  | Sequence 171, App |

## ALIGNMENTS

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: LOCATION: (1)...(625)

: OTHER INFORMATION: n = A,T,C or

US-09-825-294-210
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US-09-825-294-210
                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 210
LENGTH: 625
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Filing, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C5
                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 625; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 210, Application US/09825294 Patent No. US20020004491A1
                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                61
                                                                                                61
                                                                                                                           CAGCGACAGGCGGCAGCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGGAGGACAC 180
                                                             Conservative
                                                                                                                                                                                                           100.0%; Score 625; DB 10; 100.0%; Pred. No. 1.4e-242;
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APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Jones, Robert
ITILE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THE
TITLE OF INVENTION: COMPOSITIONS AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOSTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 10876
LENGTH: 625
CAGCGACAGGCGGCAGCACAGCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACAC
                                                                 GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
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                                                    GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF ON
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 214
LENGTH: 1897
                                                                                                                                                                                ; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(1897)
; OTHER INFORMATION: n =
US-09-825-294-214
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US-09-825-294-214
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Best Local Similarity
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                                                                                                                                Matches
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TYPE: DNA
ORGANISM: Homo
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                         CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAAGGGGAG
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Pred. No. 5.7e-235;
                                                                                                                                Mismatches
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                                                                                                                                                      Length 1897;
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RESULT 2 US-09-867-701-10876

Sequence 10876, Application Patent No. US20020132237A1

US/09867701

GENERAL INFORMATION:

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121 121 ; NAME/KEY: misc\_feature ; LOCATION: 607 ; OTHER INFORMATION: n = US-09-867-701-10876

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FEATURE:

ORGANISM: Homo sapiens

TYPE: DNA

Query Match Best Local :

Matches

al Similarity 625; Conserv

Conservative

100.0%;

.0%; Score 625; DB 10; .0%; Pred. No. 1.4e-242; 0; Mismatches 0;

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Sequence 205 Application US/09825294

Patent No. US20020004491A1

GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.

APPLICANT: Fling, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA
FILE REFERENCE: 210121.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 205

LENCTH: 1619

TYPE: DNA

ORGANISM: HOWE SERVICES
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99.8%;
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                                                                                                                                             Score 486; DB 10;
Pred. No. 9.7e-187;
0; Mismatches 0;
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                                                                                                                                                                    Length 1619;
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APPLICANT: XU, Jiangchun
APPLICANT: XU, Jiangchun
APPLICANT: XTOlk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Filng, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CF
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 211
LENGTH: 1619
TYPE: DNA
ORGANISM: Homo sapiens
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APPLICANT: Xn Tinz
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                                                                                                            CAGCGACAGGCGGCAGCACAGCACC-TGCACGAACACCGGCGAAACTGGTGCGAGGACA 179
                                                                   CCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGGA 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA 479
                                                                                        GGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GACAGTG 606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GTAAATGAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA 1471
                                                                                                                                                            606;
                                                                                                                                                                     Similarity
                                                                                                                                                            Conservative
                                                                                                                                                                     77.8%;
99.8%;
                                                                                                                                                           0,
                                                                                                                                                                    Score 486; DB 10;
Pred. No. 9.7e-187;
                                                                                                                                                          Mismatches
                                                                                                                                                                               Length 1619;
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
                                                                                                                                                                                                                                                             1 AGTTCTCCTTGCAGAGGACTGGCGCGGGGGCGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60
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                 CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
                                                                          CAGCGACAGGCGCAGCACCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC
                                                                                            CAGCGACAGGCGGCAGCACACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC
                                                                                                                                                   GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                                                                     GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OVARIAN CANCER
                                                                                                                                                                                                                                                                                                                                      Length 1362;
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APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THE TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER TITLE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT APPLICATION OUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1516
SEQ ID NO 1516
TYPE: DNA
ORGANISM: Homo sapien
US-09-867-701-1516
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                                                                                                                                               Sequence 2409, Application US/09867701 Patent No. US20020132237A1 GENERAL INFORMATION:
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APPLICANT: Aglate, Paul A.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 314;
               APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE:
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GACAGTG 1598 GACAGTG 606

US-09-825-294-208/c

GENERAL INFORMATION:

APPLICANT: Stolk, John A. APPLICANT: Algate, Paul A. APPLICANT: Fling, Steven P. APPLICANT: Xu, Jiangchun Sequence 208, Application US/09825294 Patent No. US20020004491A1

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1215

; TYPE: DNA ; ORGANISM: Homo sapiens US-09-825-294-208

SEQ ID NO 208 LENGTH: 1362

SOFTWARE: FastSEQ for Windows Version 3.0

Query Match Best Local S Matches 359

Local Similarity

Conservative

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GENERAL INFORMATION:

APPLICANT: Aglate, Paul A.

APPLICANT: Jones, Robert.

APPLICANT: Harlocker, Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.497

CURRENT APPLICATION NUMBER: US/09/867,701

CURRENT FILING DATE: 2001-05-29

NUMBER OF SEQ ID NOS: 10912

SOFTWARE: FRASTSEQ for Windows Version 4.0

SEQ ID NO 4240

LENGTH: 409
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US-09-867-701-4240
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US-09-867-701-4240/c
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LENGTH: 349
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SOFTWARE: FastSEQ for Windows Version 4.0
315
                                                                        353 ATCAGCCCTGGGCAGACACGAGCAGGAGGAGAGAAAAGAAAAAGAAAAACACAGCATGA 412
                                                                                                                                   293 GCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACA 352
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                                                                                                                  GCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACA 256
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99.78;
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                                                                                                                                                                                         Score 263; DB 10; Length 409; Pred. No. 4.5e-97;
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APPLICANT: Jones, Robert

APPLICANT: Harlocker, Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.497

CURRENT APPLICATION UNMBER: US/09/867,701

CURRENT FILING DATE: 2001-05-29

NUMBER OF SEQ ID NOS: 10912

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 8894

LENGTH: 201
                                                                 CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 1532
LENGTH: 390
                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                             Sequence 1532, Application US/09867701 Patent No. US20020132237A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity Matches 166; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                        APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
                   ORGANISM: Homo sapien FEATURE:
                                                          TYPE: DNA
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                         121 TGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAA
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100.0%; Pred. No. 4.5e-58;
n. Mismatches 0;
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; LOCATION: (1)...(558)
; OTHER INFORMATION: n = A,T,C
US-09-867-701-2375
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                                                                                                                                                                                                                                                           US-09-867-701-7826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER FILE REFERENCE: 210121.497
CURRENT APPLICATION UNMERS: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FBSESEQ for Windows Version 4.0
SEQ ID NO 2375
LENGTH: 558
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                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                               Sequence 7826, Application US/09867701 Patent No. US20020132237A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: (1)...(390)
OTHER INFORMATION: n = A,T,C or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       198 TGATGACCGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCC 257
                                                                                                                                                                                                                                                                                                                                                                                                                              134 ACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGT 193
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                                                                                                                                                                                                                                                                                                                                           194 TGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGAGCATCATGTACGCCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTAC 485
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               120;
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      PRIOR APPLICATION NUMBER: 08/346,731
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; ORGANISM: Homo sapien US-09-867-701-7826
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CURRENT APPLICATION NUMBER: US/09/938,842A
CURRENT FILING DATE: 2001-08-24
PRIOR APPLICATION NUMBER: US 60/227,866
PRIOR FILING DATE: 2000-08-24
PRIOR FILING DATE: 2000-08-24
PRIOR APPLICATION NUMBER: US 60/264,647
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-06-22
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US-09-783-590-6936
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SEQ ID NO 7826
LENGTH: 381
                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Dillon, Patrick J.
APPLICANT: Haseltine, Willia
APPLICANT: L1, Haodong
APPLICANT: Rosen, Craig A.
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                                                                                                                                                                                                                                          Sequence 6936, Application US/09783590 Patent No. US20020110850A1
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Best Local
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SEQ ID NO 1283
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Human Genes, Sequences,
FILE REFERENCE: PO-16.2C1
CURRENT APPLICATION NUMBER: US/09/783,590
CURRENT FILING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 08/420,856
PRIOR FILING DATE: 1995-04-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Wang, Xun
APPLICANT: Zhu, Tong
TITLE OF INVENTION: STRESS-REGULATED GENES OF PLANTS, TRANSGENIC PLANTS CONTAINING
TITLE OF INVENTION: SAME, AND METHODS OF USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Harper, Jeff
APPLICANT: Kreps, Joel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 819
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
nes 22; Conserv
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                                                                                                                                                                             Haseltine, William
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100.0%; Pred. No. 2.4e-28
vative 0; Mismatches (
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Sequence 698, Application US/09925299

Patent No. US20020055627A1

GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA102

CURRENT APPLICATION NUMBER: US/09/925,299

CURRENT FILING DATE: 2001-08-10

PRIOR APPLICATION NUMBER: PCT/US00/05883

PRIOR APPLICATION NUMBER: 60/124,270

PRIOR APPLICATION NUMBER: 60/124,270

PRIOR FILING DATE: 1999-03-12

NUMBER OF SEQ ID NOS: 1556
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Best Local :
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NUMBER OF SEQ ID NOS: 12485
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 6936
LENGTH: 322
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LOCATION: (292)
OTHER INFORMATION: n equals a,t,g,
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LOCATION: (241)
OTHER INFORMATION: n equals a,t,g,
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LOCATION: (236)
OTHER INFORMATION: n equals a,t,g, or c
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LOCATION: (216)
OTHER INFORMATION: n equals a,t,g,
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LOCATION: (258)
OTHER INFORMATION: n equals a,t,g, or c
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LOCATION: (20)
OTHER INFORMATION: n equals a,t,g,
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100.0%; Pred. No. 8.4;
ative 0; Mismatches
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                                                                                                                           ; OTHER INFORMATION: n = A,T,C or G US-09-834-975-379
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                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 379 LENGTH: 547
                                                            Best Local
Matches
                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 379, Application US/09834975 Patent No. US20020110815A1
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/834,975
CURRENT FILING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: 60/197,538
PRIOR FILING DATE: 2000-04-14
NUMBER OF SEQ ID NOS: 1046
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                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: BOOM, Jeffrey
APPLICANT: BOIL, ANDREW
APPLICANT: WON HUMSEN, COMPOSITIONS AND METHODS
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE IDENTIFICATION, ASSESSMENT, PREVENTION, AND THERAPY
TITLE OF INVENTION: OF HUMAN CANCERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Lillie, James APPLICANT: Brown, Jeffre
                                                                                                                                                    NAME/KEY: misc_feature LOCATION: (1)...(547)
                                                                                                                                                                                                  ORGANISM: Homo sapiens
                                                                                                                                                                                                                        TYPE: DNA
                                                                                                                                                                                           FEATURE:
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LOCATION: (499)
OTHER INFORMATION: n equals a,t,g,
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LOCATION: (496)
OTHER INFORMATION: n equals a,t,g,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature
LOCATION: (480)
OTHER INFORMATION: n equals a,t,g,
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NAME/KEY: misc_feature
LOCATION: (449)
OTHER INFORMATION: n equals a,t,g,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
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               605 TGNAAAAAAAAAAAAAAAAA 625
527 TGNAAAAAAAAAAAAAAAA 547
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mes 21; Conserv
                                                                            Similarity
                                                            Conservative
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100.0%; F1
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Pred. No.
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US-09-783-590-7037
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OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (109)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (122)
OTHER INFORMATION: n equals a,t,g, or c
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PRIOR APPLICATION NUMBER: 08/346,731
PRIOR FILING DATE: 1994-11-21
NUMBER OF SEQ ID NOS: 12485
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 7037
LENGTH: 600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Human Genes, Sequences, and Expression Products 16.2 FILE REFERENCE: PO-16.2C1
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OTHER INFORMATION: n equals a,t,
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OTHER INFORMATION: n equals a,t,g,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
                                                                                                                          NAME/KEY: misc feature
LOCATION: (409)
OTHER INFORMATION: n equals a,t,g, or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: n equals a,t,g, or
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OTHER INFORMATION: n equals a,t,g, or
                                                              LOCATION: (418)
OTHER INFORMATION: n equals
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                        NAME/KEY: misc feature LOCATION: (428)
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US-09-925-301-539
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                                                                                                                                                                                                SOFTWARE: PatentIn Ver. SEQ ID NO 539
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                                                                                                                                                                                                                                                                                                      APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins
FILE REFERENCE: PA106
CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/124,270 PRIOR FILING DATE: 1999-03-12 NUMBER OF SEQ ID NOS: 1694
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OTHER INFORMATION: n equals a,t,g, or NAME, KEY: misc feature
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OTHER INFORMATION: n equals a,t,g, or NAME/KEY: misc feature
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OTHER INFORMATION: n equals a,t,g, or
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OTHER INFORMATION: n equals a,t,g, or
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      LOCATION: (1305)
OTHER IFFORMATION: n equals a,t,g, or NAME/KBY: misc_feature
LOCATION: (1344)
                                                                                                                                   TYPE: DNA
ORGANISM: Homo sapiens
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NAME/KEY: misc_feature
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es 21; Conserv
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Pred. No.
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Indels

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Gaps

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and Antibodies

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CURRENT APPLICATION NUMBER: US/09/925,297
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05989
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 928
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 171
APPLICANT: Rosen, Craig A.

APPLICANT: Ruben, Steven M.

TITLE OF INVENTION: Human Genes, Sequences, an FILE REFERENCE: PO-16.201

CURRENT APPLICATION NUMBER: US/09/783,590

CURRENT FILING DATE: 2000-02-15

PRIOR APPLICATION NUMBER: 08/420,856

PRIOR FILING DATE: 1995-04-12

PRIOR FILING DATE: 1995-04-12

PRIOR FILING DATE: 1994-11-21

PRIOR APPLICATION NUMBER: 08/346,731

PRIOR FILING DATE: 1994-11-21

NUMBER OF SEQ ID NOS: 12485

SOFTWARE: Patentin Ver. 2.0
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US-09-925-297-171
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Best Local Similarity 100.0%; P
Watches 21; Conservative 0;
                                                                                                                                                                                                                                                                                                                                       Sequence 2406, Application US/09783590 Patent No. US20020110850A1
                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                APPLICANT: Dillon, Patrick J.
APPLICANT: Haseltine, Willis
APPLICANT: Li, Haodong
APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
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TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA105
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LOCATION: (1370)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: misc feature
LOCATION: (1400)
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OTHER INFORMATION: n equals a,t,g, or c
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les 21; Conserv
                                                                                                                                                                                                                                                                          Haseltine, William A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.4%; Score 21; DB 10; Length 1402; 100.0%; Pred. No. 7.6;
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Pred. No.
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                                                                                                                                                                                    and Expression Products 16.2
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RESULT 23
US-09-925-301-759/c
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Sequence 759, Application US/09925301
Patent No. US20020052308A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA106
                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 934

SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 419
LENGTH: 60
                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Pyle, Ruth
APPLICANT: Xu, Jiangchun
APPLICANT: Secrist, Heather
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.552
CURRENT APPLICATION NUMBER: US/09/919,580
CURRENT FILING DATE: 2001-07-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 419, Application US/09919580 Patent No. US20020110832A1
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LENGTH: 58
                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature LOCATION: 54
                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc feature
LOCATION: (58)
OTHER INFORMATION: n equals a,t,g, or c
                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc feature
LOCATION: (52)
OTHER INFORMATION: n equals a,t,g,
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LOCATION: (34)
OTHER INFORMATION: n equals a,t,g, or
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                                                                                                                                                                                        606 GNAAAAAAAAAAAAAAAA 625
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                                                                                                                                                                      55 GNAAAAAAAAAAAAAAA 36
                                                                                                                                                                                                                                                          Local Similarity
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                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                      100.0%;
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100.0%; Pred. No.
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100.0%; Pred. No.
                                                                                                                                                                                                                                        0;
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23;
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                                                                                                                                                                                                                                                                    10; Length 60;
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                                                                                                                                                                                                                                    Indels
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; NAME/KEY: misc_feature; LOCATION: (66); OTHER INFORMATION: n equals a,t,g, or c US-09-925-301-759
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PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTMARE: Patentin Ver. 2.0
SEQ ID NO 759
LENGTH: 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 241, Application US/09815343 Patent No. US20010055596A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
                                                                                                                                                                                                                                                                                                                                              APPLICANT: Meagher, Madeleine
APPLICANT: Xu, Jiangchun
APPLICANT: Xiu, Jiangchun
APPLICANT: King, Gordon E.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121-504
CURRENT APPLICATION NUMBER: US/09/815,343
CURRENT FILING DATE: 2001-03-22
                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 1556
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: n equals a,t,g, or NAME/KEY: misc_feature LOCATION: (63)
OTHER INFORMATION: n equals a,t,g, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 20; Pred. No.
                                                                                    Score 20; Pred. No.
                                                                    Mismatches
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. 23;
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23;
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APPLICANT: Wang, Ton
APPLICANT: MCNeill;
APPLICANT: Watanabe
APPLICANT: Carter,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-895-828-248
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US-10-046-935-291/c
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                                                                                                                                              US-10-046-935-291
                                                                                                                                                                                                                                                                          TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.527C1
CURRENT APPLICATION NUMBER: US/10/046,935
CURRENT FILING DATE: 2002-01-15
NUMBER OF SEQ ID NOS: 2239
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 291
LENGTH: 226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 248
LENGTH: 217
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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Best Local Similarity
Matches 20; Conserv
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                                                                          Matches
                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Henderson, Robert A.
APPLICANT: Kalos, Michael D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Jiang, Yuqiu
APPLICANT: Harlocker, Susan
APPLICANT: Secrist, Heather
APPLICANT: Wang, Aijun
APPLICANT: Stolk, John A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/895,828
CURRENT FILING DATE: 2001-06-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
                                                                                                                                                                  NAME/KEY: misc_feature
LOCATION: 20, 57, 141, 184, 206
OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                         ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                              TYPE: DNA
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                                   606 GNAAAAAAAAAAAAAAAA 625
 21 GNAAAAAAAAAAAAAAAAAA
                                                                                      Local Similarity
                                                                          20;
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Carter, Darrick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        McNeill, Patricia D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                        Conservative
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, 100.0%; Prr
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Pred. No.
                                                                            Mismatches
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. 22;
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22;
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                                                                                                                 Length 226;
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606 GNAAAAAAAAAAAAAAA 625

60 GNAAAAAAAAAAAAAAAA 41

Query Match Best Local 9

Local Similarity

100.0%;

3.2%;

NAME/KEY: misc\_feature
LOCATION: (65)
OTHER INFORMATION: n equals a,t,g,

FEATURE:

NAME/KEY: misc\_feature

LOCATION: (6)
OTHER INFORMATION: n equals a,t,g,

NAME/KEY: misc\_feature

LOCATION: (59)

ORGANISM: Homo sapiens

TYPE: DNA

Matches

20;

Conservative

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RESULT 24 US-09-815-343-241/c

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606 GNAAAAAAAAAAAAAA 625

88 GNAAAAAAAAAAAAAAAA 69

Matches Query Match

Best Local

Similarity

100.0%;

3.28;

Conservative

0,

US-09-815-343-241

NAME/KEY: misc\_feature LOCATION: (1)...(97) OTHER INFORMATION: n = A,T,C or G

SEQ ID NO 241 LENGTH: 97

ENGTH:

TYPE: DNA
ORGANISM: Homo sapien
FEATURE:

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US-09-834-975-374/c
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; Sequence 46, Application US/10040916
; Patent No. US20020146769A1
; GENERAL INFORMATION:
                                                                                                                     Sequence 374, Application US/09834975 Patent No. US20020110815A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
APPLICANT: Lillie, James
APPLICANT: Brown, Jeffrey
APPLICANT: Bott, Andrew
APPLICANT: Wan Huffel, Christophe
APPLICANT: Van Huffel, Christophe
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS AND METHOD:
TITLE OF INVENTION: FOR THE IDENTIFICATION, ASSESSMENT,
                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
TELECOMMUNICATION: INFORMATION:
TELEPHONE: (617) 498-8224
TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 46:
                                                                                                                                                                                                                                                                                  606 GNAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                                                                                           245 GNAAAAAAAAAAAAAA 264
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 08/686,878 FILING DATE: <Unknown> ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/887,029
FILING DATE: 07-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE DESCRIPTION: SEQ ID NO: 46:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute,
STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Jacobs, Kenneth McCoy, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/10/040,916 FILING DATE: 07-Jan-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Cambridge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 265 base pairs
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                                                                                                                                                                                                                                                                                                                                          Conservative
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Evans, Cheryl
Spaulding, Vikki
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LaVallie, Edward
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Racie, Lisa
                                                                                                                                                                                                                                                                                                                                                         100.0%;
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100.0%; Pred. No.
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                      COMPOSITIONS AND METHODS
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                                                                                                                                                                                                                                                                                                                                                                           Length 265;
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  PREVENTION, AND THERAPY
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                                                                                                                                                                                                                                                                                                                                    Gaps
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US-09-783-590-10830
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US-09-777-564-625/c
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; NAME/KEY: misc_feature
; LOCATION: (1)...(291)
; OTHER INFORMATION: n = A,T,C or
US-09-834-975-374
                                                                                                                                                                         Sequence 10830, Application US/09783590 Patent No. US20020110850A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Algate, Paul A.
APPLICANT: Mannion, Jane
APPLICANT: Mannion, Jane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/777,564
CURRENT FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 1730
SOFTWARE: FastSEQ for Window Version 4.0
SEQ ID NO 625
LENGTH: 451
                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 625, Application US/09777564
Patent No. US20020022591A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 1046
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 374
LENGTH: 291
TYPE: DNA
          APPLICANT: Rosen, Craiq A.
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Human Genes, Sequences, and Expression Products 16.2
FILE REFERENCE: PO-16.2C1
                                                                                             APPLICANT: Dillon, Patrick J.
APPLICANT: Haseltine, William A.
APPLICANT: Li, Haodong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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CURRENT APPLICATION NUMBER: US/09/783,590
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER FILE REFERENCE: 210121.493
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/197,538 PRIOR FILING DATE: 2000-04-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/834,975
CURRENT FILING DATE: 2001-04-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: n = A,T,C or G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature LOCATION: (1)...(451)
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100.0%; Pred. No.
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SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 10830
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PRIOR FILING DATE: 1995-04-12
PRIOR APPLICATION NUMBER: 08/346,731
PRIOR FILING DATE: 1994-11-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 12485
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                                                                         LOCATION: (392)
OTHER INFORMATION: n equals a,t,g, or
NAME/KEY: misc feature
LOCATION: (396)
OTHER INFORMATION: n equals a,t,g, or
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OTHER INFORMATION: n equals a,t,g, or
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OTHER INFORMATION: n equals a,t,g,
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OTHER INFORMATION: n equals a,t,g,
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                    OTHER INFORMATION: n equals a,t,g, or c
                                     NAME/KEY: misc feature LOCATION: (404)
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misc feature
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PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 819
LENGTH: 648
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No. US20020052308A1 GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/925,301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies FILE REFERENCE: PA106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Rosen et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc feature
LOCATION: (419)
OTHER INFORMATION: n equals a,t,9,
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OTHER INFORMATION: n equals
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OTHER INFORMATION: n
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             OTHER INFORMATION: n equals a,t,g,
NAME/KEY: misc_feature
LOCATION: (584)
                                                                   OTHER INFORMATION: n equals a,t,g, or NAME/KEY: misc_feature LOCATION: (565)
                                                                                                                         OTHER INFORMATION: n equals a,t,g, NAME/KEY: misc_feature LOCATION: (547)
                                                                                                                                                                                                 LOCATION: (518)
OTHER INFORMATION: n equals
NAME/KEY: misc_feature
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LOCATION: (369)
OTHER INFORMATION: n equals a,t,9, or
                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
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OTHER INFORMATION: n equals a,t,g, or
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APPLICANT: Wang, Aijun
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS
TITLE OF INVENTION: COMPOSITIONS AND METHODS
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CA
FILLE REFERENCE: 21021, 427C23
CURRENT APPLICATION NUMBER: US/09/759,143
CURRENT FILING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 934
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                 US-09-780-669-32/c
                                                                                                                                                    RESULT 33
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                                                                                        Sequence 32, Application US/09780669 Patent No. US20020051977A1 GENERAL INFORMATION:
                                             APPLICANT:
APPLICANT:
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LOCATION: (1)...(789)
THER INFORMATION: n = A,T,C or G
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mes 20; Conserv
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                                   Xu, Jiangchun
Dillon, Davin C.
Mitcham, Jennifer L.
Jiang, Yuqui
Henderson, Robert A.
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Vedvick,
                              Harlocker, Susan L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Carter, Darrick
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                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPOSITIONS AND METHODS FOR THE THERAPY AND DIAGNOSIS OF PROSTATE CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.08; P1
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Thomas S.
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Pred. No.
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RESULT 34
US-09-030-606-32/c
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Patent No. US20020081580A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Xi, Jiangchun
APPLICANT: Dillon, Davin C.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNODIAGNOSIS OF PROSTATE CANCER AND METHOD
NUMBER OF SEQUENCES: 224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best
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CURRENT FILING DATE: 2001-02-09
NUMBER OF SEQ ID NOS: 943
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
LENGTH: 789
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INFORMATION FOR SEQ ID NO:
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APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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TYPE: DNA
ORGANISM: Homo sapien
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LOCATION: (1)...(789)
OTHER INFORMATION: n = A,T,C or G
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                                                                    CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
NAME: Maki, David J.
31,392
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                        REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                            SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/030,606
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                CITY: Seattle
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les 20; Conserv
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Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
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Hepler, William
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Carter, Darrick
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                 (206)
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               682-6031
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                                                               210121.428C3
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20;
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US-09-822-827-32/c
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US-09-115-453-32/c
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US-09-115-453-32
                                                                                                                                                                                                                                                                                                                      Sequence 32, Application US/09115453B Patent No. US20020090372A1 GENERAL INFORMATION:
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Patent NO. US20020081680A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
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                                                                                                                                         NUMBER OF SEQ ID NOS: 228
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 32
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CURRENT FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER EILE REFERENCE: 210121.534C1
                                                                                                                                                                                             TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY OF PROSTATE CANCER AND TITLE OF INVENTION: METHODS FOR THEIR USE FILE REFERENCE: 210121.427C4
CURRENT APPLICATION NUMBER: US/09/115,453B
CURRENT FILING DATE: 1998-07-14
                                                                                                                                                                                                                                                                                      APPLICANT: Xu, Jiangchun APPLICANT: Dillon, Davin C.
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         FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(789)
OTHER INFORMATION: n = A,T,C or G
                                                                              LENGTH: 789
TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: n = A,T,C or G
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LENGTH: 789 base pairs
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Pred. No.
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US-09-783-590-8086/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 3, Application US/09949654 Patent No. US20020127644A1
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Best Local Similarity
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                                                                                                                                                       SEQ ID NO 8086
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/949,654
CURRENT FILING DATE: 2001-09-12
PRIOR APPLICATION NUMBER: 60/231,572
PRIOR FILING DATE: 2000-09-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: CL000817
                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Dillon, Patrick J.
APPLICANT: Haseltine, William A.
APPLICANT: Li, Haodong
APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
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                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/783,590
CURRENT FILING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 08/420,856
PRIOR FILING DATE: 1995-04-12
PRIOR APPLICATION NUMBER: 08/346,731
PRIOR FILING DATE: 1994-11-21
NUMBER OF SEQ ID NOS: 12485
                                                                                                                                                                                                                                                                                                                                 APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Human Genes, Sequences,
FILE REFERENCE: PO-16.2C1
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                                                                                                                                                                         SOFTWARE: PatentIn Ver.
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ORGANISM: Human
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              FEATURE: misc feature
NAME/KEY: misc feature
LOCATION: (10)
OTHER INFORMATION: n equals a,t,g, or c
                                                                                             LENGTH: 51
TYPE: DNA
ORGANISM: Homo sapiens
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NAME/KEY: misc feature
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US-09-919-580-705/c
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                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/919,580
CURRENT FILING DATE: 2001-07-30
NUMBER OF SEQ ID NOS: 934
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 705
LENGTH: 53
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APPLICANT: Xu, Jiangchun
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
ETLE REFERENCE: 210121.552
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SEQ ID NO 5881
LENGTH: 52
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TITLE OF INVENTION: Human Genes, Sequences, and Expression Products 16.2

EILE REFERENCE: PO-16, 201

CURRENT APPLICATION NUMBER: US/09/783,590

CURRENT FILING DATE: 2000-02-15

PRIOR APPLICATION NUMBER: 08/420,856

PRIOR FILING DATE: 1995-04-12

PRIOR FILING DATE: 1995-04-12

PRIOR APPLICATION NUMBER: 08/346,731

PRIOR PRIOR PRIOR APPLICATION NUMBER: 08/346,731

PRIOR PRIOR PRIOR PRIOR PRIOR E: 1994-11-21

PRIOR PRIOR PRIOR DATE: 1995-04-12

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                                                              ORGANISM: Homo sapiens
                                                                                                           TYPE: DNA
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NAME/KEY: misc_feature
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LOCATION: (48)
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ORGANISM: Homo sapiens
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OTHER INFORMATION: n equals a,t,g, or c
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Ruben, Steven M.
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100.0%; Pred. No.
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CURRENT APPLICATION NUMBER: US/09/783,590
CURRENT ETLING DATE: 2000-02-15
PRIOR APPLICATION NUMBER: 08/420,856
PRIOR FILING DATE: 1995-04-12
PRIOR APPLICATION NUMBER: 08/346,731
PRIOR FILING DATE: 1994-11-21
NUMBER OF SEQ ID NOS: 12485
SECTWARE: PATENTIN VET. 2.0
SEQ ID NO 3220
LENGTH: 53
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US-09-783-590-3220/c
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                 NUMBER OF SEQ ID NOS: 934

SOFTWARE: FRASTSEQ for Windows Version 4.0
SEQ ID NO 171
LENGTH: 56
                                                                                                                                                                                                                                                     Sequence 171, Application US/09919580 Patent No. US20020110832A1
                                                                                                       APPLICANT: Pyle, Ruth
APPLICANT: Xu, Jiangchun
APPLICANT: Su, Jiangchun
APPLICANT: Su, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.552
                                                             CURRENT APPLICATION NUMBER: US/09/919,580 CURRENT FILING DATE: 2001-07-30 NUMBER OF SEQ ID NOS: 934
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APPLICANT: Dillon, Patrick J.
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TYPE: DNA
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APPLICANT: Li, Haodong
APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steven M.
TITLE OF INVENTION: Human Genes, Sequences, and Expression Products 16.2
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OTHER INFORMATION: n equals a,t,g, or c
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ORGANISM: Homo sapiens
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OTHER INFORMATION:
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les 19; Conserva
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100.0%; Pred. No.
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US-09-919-580-713/c
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SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 677
LENGTH: 56
TYPEF: 5...
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                                                                                                           NUMBER OF SEQ ID NOS: 934
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 713
LENGTH: 56
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APPLICANT: Xu, Jiangehun
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.552
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/919,580 CURRENT FILING DATE: 2001-07-30
                                                                                                                                                                                        APPLICANT: Pyle, Ruth
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.552
CURRENT APPLICATION NUMBER: US/09/919,580
CURRENT FILING DATE: 2001-07-30
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LOCATION: 49
OTHER INFORMATION: n = A,T,C or G
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LOCATION: 53
OTHER INFORMATION: n = A,T,C or G
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TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: 35, 37, 42, 56
OTHER INFORMATION: n = A,T,C or G
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100.0%; pred. No.
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US-09-925-301-758/c
                                                                          Db
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Search completed: November 8, 2002, 02:06:25 Job time : 100.162 secs
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SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 758
LEMOTOR
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Best Local Similarity
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PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
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                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
LOCATION: (45)
OTHER INFORMATION: n equals a,t,g, or
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LOCATION: (38)
OTHER INFORMATION: n equals a,t,g, or
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LOCATION: (36)
OTHER INFORMATION: n equals a,t,g, or c
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LOCATION: (40)
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                                                                                                                                                                Local Similarity
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Minimum DB seq length: 0
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Copyright (c) 1993 - 2002 Compugen Ltd.
/cgn2_6/ptodata/1/pna/US102B_COMB.seq:*
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| 4: //cgn2_6/ptodata/1/pna/US6000_COMB. 6: //cgn2_6/ptodata/1/pna/US60002_COMB. 6: //cgn2_6/ptodata/1/pna/US6002_COMB. 7: //cgn2_6/ptodata/1/pna/US6003_COMB. 7: //cgn2_6/ptodata/1/pna/US6003_COMB. 9: //cgn2_6/ptodata/1/pna/US6006_COMB. 1: //cgn2_6/ptodata/1/pna/US6009_COMB. 2: //cgn2_6/ptodata/1/pna/US6009_COMB. 2: //cgn2_6/ptodata/1/pna/US6011_COMB. 3: //cgn2_6/ptodata/1/pna/US6011_COMB. 4: //cgn2_6/ptodata/1/pna/US6013_COMB. 5: //cgn2_6/ptodata/1/pna/US6013_COMB. 6: //cgn2_6/ptodata/1/pna/US6013_COMB. 7: //cgn2_6/ptodata/1/pna/US6013_COMB. 7: //cgn2_6/ptodata/1/pna/US6013_COMB. 7: //cgn2_6/ptodata/1/pna/US6019_COMB. 9: //cgn2_6/ptodata/1/pna/US6019_COMB. 1: //cgn2_6/ptodata/1/pna/US6019_COMB. 1: //cgn2_6/ptodata/1/pna/US6020_COMB. 2: //cgn2_6/ptodata/1/pna/US6020_COMB. 3: //cgn2_6/ptodata/1/pna/US6023_COMB. 3: //cgn2_6/ptodata/1/pna/US6023_COMB. 3: //cgn2_6/ptodata/1/pna/US6023_COMB. 3: //cgn2_6/ptodata/1/pna/US6023_COMB. 3: //cgn2_6/ptodata/1/pna/US6023_COMB. 3: //cgn2_6/ptodata/1/pna/US6023_COMB. 3: //cgn2_6/ptodata/1/pna/US6033_COMB. 4: //cgn2_6/ptoda | gn2_6/ptodata/1/pna/US6041_COMB.             | 85:         |  |
|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------|-------------|--|
| 4: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6004_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6010_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6011_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6022_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6022_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6022_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6032_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6032_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6032_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6033_COMB.s. n2_6/ptodata/1/pna/US6040 COMB.             | 84:         |  |
| 4: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6004_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 1: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 1: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 1: //cgn2_6/ptodata/1/pna/US6011_COMB.s. 1: //cgn2_6/ptodata/1/pna/US6011_COMB.s. 1: //cgn2_6/ptodata/1/pna/US6011_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6011_COMB.s. 1: //cgn2_6/ptodata/1/pna/US6011_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6011_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6011_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6011_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6011_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6033_COMB.s.                                                                                                                                                                  | gn2_6/ptodata/1/pna/US6039_COMB.             | 83:         |  |
| 4: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s. 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60006_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60010_COMB.s. 9: //cgn2_6/ptodata/1/pna/US601010_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60112_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6022_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6033_COMB.s.                                                                                                                                                                                               | <pre>gn2_6/ptodata/1/pna/US6038_COMB.</pre>  | 82:         |  |
| 4: //cgn2_6/ptodata/1/pna/US6000_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6004_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s. 1: //cgn2_6/ptodata/1/pna/US60007_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60010_COMB.s. 1: //cgn2_6/ptodata/1/pna/US6010_COMB.s. 3: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 5: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 6: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 6: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 8: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6013_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6022_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6033_COMB.s.                                                                                                                                                                                                                                                                                                                                                                                                           | gn2_6/ptodata/1/pna/US6037_COMB.s            | 81:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: /cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: /cgn2_6/ptodata/1/pna/US6001_COMB.s. 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s. 0: /cgn2_6/ptodata/1/pna/US6006_COMB.s. 0: /cgn2_6/ptodata/1/pna/US6006_COMB.s. 0: /cgn2_6/ptodata/1/pna/US6001_COMB.s. 0: /cgn2_6/ptodata/1/pna/US6001_COMB.s. 0: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6012_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6022_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6022_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6025_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6025_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6025_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6025_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6025_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6032_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6032_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6033_COMB.s.                                                                                                                                                                                                                                                                                                                                                                                                                                                                | <pre>gn2_6/ptodata/1/pna/US6036_COMB.s</pre> | 80:         |  |
| 4: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s. 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s. 0: //cgn2_6/ptodata/1/pna/US60006_COMB.s. 2: //cgn2_6/ptodata/1/pna/US60010_COMB.s. 3: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 5: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 5: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 6: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 6: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6033_COMB.s.                                                                                                                                                                                                                                        | gn2_6/ptodata/1/pna/US6035_COMB.s            | 79:         |  |
| 4: //cgn2_6/ptodata/1/pna/US6001_COMB.s<br>6: //cgn2_6/ptodata/1/pna/US6001_COMB.s<br>7: //cgn2_6/ptodata/1/pna/US6003_COMB.s<br>7: //cgn2_6/ptodata/1/pna/US6003_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6006_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6006_COMB.s<br>0: //cgn2_6/ptodata/1/pna/US60010_COMB.s<br>0: //cgn2_6/ptodata/1/pna/US6010_COMB.s<br>0: //cgn2_6/ptodata/1/pna/US60110_COMB.s<br>3: //cgn2_6/ptodata/1/pna/US60112_COMB.s<br>5: //cgn2_6/ptodata/1/pna/US60112_COMB.s<br>6: //cgn2_6/ptodata/1/pna/US60113_COMB.s<br>6: //cgn2_6/ptodata/1/pna/US60113_COMB.s<br>6: //cgn2_6/ptodata/1/pna/US60113_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US60112_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US60112_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US60112_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6012_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6012_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6012_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6012_COMB.s<br>1: //cgn2_6/ptodata/1/pna/US6022_COMB.s<br>2: //cgn2_6/ptodata/1/pna/US6023_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6023_COMB.s<br>1: //cgn2_6/ptodata/1/pna/US6023_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6023_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6023_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6023_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6023_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6023_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6023_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6033_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6033_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6033_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6033_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6033_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             | gn2_6/ptodata/1/pna/US6034_COMB.s            | 78:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6001_COMB.s 5: /cgn2_6/ptodata/1/pna/US6001_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 1: /cgn2_6/ptodata/1/pna/US6001_COMB.s 2: /cgn2_6/ptodata/1/pna/US6011_COMB.s 3: /cgn2_6/ptodata/1/pna/US6011_COMB.s 5: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 7: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6012_COMB.s 9: /cgn2_6/ptodata/1/pna/US6013_COMB.s 9: /cgn2_6/ptodata/1/pna/US6013_COMB.s 9: /cgn2_6/ptodata/1/pna/US6013_COMB.s 9: /cgn2_6/ptodata/1/pna/US6013_COMB.s 9: /cgn2_6/ptodata/1/pna/US6021_COMB.s 9: /cgn2_6/ptodata/1/pna/US6022_COMB.s 9: /cgn2_6/ptodata/1/pna/US6023_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             | gn2_6/ptodata/1/pna/US6033_COMB.s            | 77:         |  |
| 4: //cgn2_6/ptodata/1/pna/US6001_COMB.s 5: //cgn2_6/ptodata/1/pna/US6001_COMB.s 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s 0: //cgn2_6/ptodata/1/pna/US6006_COMB.s 2: //cgn2_6/ptodata/1/pna/US60010_COMB.s 3: //cgn2_6/ptodata/1/pna/US6010_COMB.s 5: //cgn2_6/ptodata/1/pna/US6011_COMB.s 6: //cgn2_6/ptodata/1/pna/US6011_COMB.s 6: //cgn2_6/ptodata/1/pna/US6011_COMB.s 6: //cgn2_6/ptodata/1/pna/US6011_COMB.s 9: //cgn2_6/ptodata/1/pna/US6011_COMB.s 9: //cgn2_6/ptodata/1/pna/US6011_COMB.s 9: //cgn2_6/ptodata/1/pna/US6011_COMB.s 9: //cgn2_6/ptodata/1/pna/US6011_COMB.s 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s 9: //cgn2_6/ptodata/1/pna/US6022_COMB.s 9: //cgn2_6/ptodata/1/pna/US6023_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | gn2_6/ptodata/1/pna/US6032_COMB.s            | 76:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s 6: /cgn2_6/ptodata/1/pna/US6001_COMB.s 7: /cgn2_6/ptodata/1/pna/US6001_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US60006_COMB.s 0: /cgn2_6/ptodata/1/pna/US60010_COMB.s 0: /cgn2_6/ptodata/1/pna/US6010_COMB.s 3: /cgn2_6/ptodata/1/pna/US60110_COMB.s 5: /cgn2_6/ptodata/1/pna/US60113_COMB.s 6: /cgn2_6/ptodata/1/pna/US60113_COMB.s 6: /cgn2_6/ptodata/1/pna/US60113_COMB.s 6: /cgn2_6/ptodata/1/pna/US60113_COMB.s 6: /cgn2_6/ptodata/1/pna/US60113_COMB.s 9: /cgn2_6/ptodata/1/pna/US60113_COMB.s 9: /cgn2_6/ptodata/1/pna/US60113_COMB.s 9: /cgn2_6/ptodata/1/pna/US60113_COMB.s 9: /cgn2_6/ptodata/1/pna/US6012_COMB.s 9: /cgn2_6/ptodata/1/pna/US6013_COMB.s 9: /cgn2_6/ptodata/1/pna/US6013_COMB.s 9: /cgn2_6/ptodata/1/pna/US6013_COMB.s 9: /cgn2_6/ptodata/1/pna/US6022_COMB.s 9: /cgn2_6/ptodata/1/pna/US6023_COMB.s 9: /cgn2_6/ptodata/1/pna/US6023_COMB.s 9: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  | <pre>gn2_6/ptodata/1/pna/US6031_COMB.s</pre> | 75:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6001_COMB.s 6: /cgn2_6/ptodata/1/pna/US6001_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6004_COMB.s 8: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US6001_COMB.s 1: /cgn2_6/ptodata/1/pna/US6001_COMB.s 3: /cgn2_6/ptodata/1/pna/US6011_COMB.s 5: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 7: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6012_COMB.s 1: /cgn2_6/ptodata/1/pna/US6013_COMB.s 1: /cgn2_6/ptodata/1/pna/US6020_COMB.s 1: /cgn2_6/ptodata/1/pna/US6020_COMB.s 1: /cgn2_6/ptodata/1/pna/US6023_COMB.s 1: /cgn2_6/ptodata/1/pna/US6024_COMB.s 1: /cgn2_6/ptodata/1/pna/US6024_COMB.s 1: /cgn2_6/ptodata/1/pna/US6023_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | <pre>gn2_6/ptodata/1/pna/US6030_COMB.s</pre> | 74:         |  |
| 4: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s. 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s. 0: //cgn2_6/ptodata/1/pna/US60006_COMB.s. 0: //cgn2_6/ptodata/1/pna/US60010_COMB.s. 2: //cgn2_6/ptodata/1/pna/US6010_COMB.s. 3: //cgn2_6/ptodata/1/pna/US60112_COMB.s. 5: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 6: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 6: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 6: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 7: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 1: //cgn2_6/ptodata/1/pna/US6013_COMB.s. 1: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 5: //cgn2_6/ptodata/1/pna/US6022_COMB.s. 5: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6023_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6026_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6026_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6026_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6026_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6026_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6026_COMB.s.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     | <pre>gn2_6/ptodata/1/pna/US6029_COMB.s</pre> | 73:         |  |
| 4: //cgn2_6/ptodata/1/pna/US6001_COMB.s 6: //cgn2_6/ptodata/1/pna/US6001_COMB.s 7: //cgn2_6/ptodata/1/pna/US6002_COMB.s 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s 0: //cgn2_6/ptodata/1/pna/US6006_COMB.s 0: //cgn2_6/ptodata/1/pna/US60010_COMB.s 0: //cgn2_6/ptodata/1/pna/US6010_COMB.s 3: //cgn2_6/ptodata/1/pna/US6011_COMB.s 5: //cgn2_6/ptodata/1/pna/US6011_COMB.s 6: //cgn2_6/ptodata/1/pna/US6011_COMB.s 6: //cgn2_6/ptodata/1/pna/US6011_COMB.s 6: //cgn2_6/ptodata/1/pna/US6011_COMB.s 9: //cgn2_6/ptodata/1/pna/US6011_COMB.s 9: //cgn2_6/ptodata/1/pna/US6011_COMB.s 9: //cgn2_6/ptodata/1/pna/US6011_COMB.s 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s 9: //cgn2_6/ptodata/1/pna/US6013_COMB.s 9: //cgn2_6/ptodata/1/pna/US6013_COMB.s 9: //cgn2_6/ptodata/1/pna/US6013_COMB.s 9: //cgn2_6/ptodata/1/pna/US6022_COMB.s 9: //cgn2_6/ptodata/1/pna/US6022_COMB.s 9: //cgn2_6/ptodata/1/pna/US6022_COMB.s 9: //cgn2_6/ptodata/1/pna/US6025_COMB.s 9: //cgn2_6/ptodata/1/pna/US6025_COMB.s 9: //cgn2_6/ptodata/1/pna/US6025_COMB.s 9: //cgn2_6/ptodata/1/pna/US6025_COMB.s 9: //cgn2_6/ptodata/1/pna/US6025_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | <pre>gn2_6/ptodata/1/pna/US6028_COMB.s</pre> | 72:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s 5: /cgn2_6/ptodata/1/pna/US6001_COMB.s 6: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US6001_COMB.s 0: /cgn2_6/ptodata/1/pna/US6001_COMB.s 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s 3: /cgn2_6/ptodata/1/pna/US6011_COMB.s 5: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 0: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 0: /cgn2_6/ptodata/1/pna/US6011_COMB.s 0: /cgn2_6/ptodata/1/pna/US6012_COMB.s 0: /cgn2_6/ptodata/1/pna/US6013_COMB.s 0: /cgn2_6/ptodata/1/pna/US6013_COMB.s 0: /cgn2_6/ptodata/1/pna/US6013_COMB.s 0: /cgn2_6/ptodata/1/pna/US6022_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6023_COMB.s 0: /cgn2_6/ptodata/1/pna/US6024_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            | <pre>gn2_6/ptodata/1/pna/US6027_COMB.s</pre> | 71:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: /cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: /cgn2_6/ptodata/1/pna/US6003_COMB.s. 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s. 0: /cgn2_6/ptodata/1/pna/US6006_COMB.s. 0: /cgn2_6/ptodata/1/pna/US60009_COMB.s. 2: /cgn2_6/ptodata/1/pna/US60010_COMB.s. 2: /cgn2_6/ptodata/1/pna/US60110_COMB.s. 3: /cgn2_6/ptodata/1/pna/US60110_COMB.s. 5: /cgn2_6/ptodata/1/pna/US60110_COMB.s. 6: /cgn2_6/ptodata/1/pna/US60110_COMB.s. 6: /cgn2_6/ptodata/1/pna/US60110_COMB.s. 6: /cgn2_6/ptodata/1/pna/US60110_COMB.s. 9: /cgn2_6/ptodata/1/pna/US60110_COMB.s. 9: /cgn2_6/ptodata/1/pna/US60110_COMB.s. 9: /cgn2_6/ptodata/1/pna/US60110_COMB.s. 9: /cgn2_6/ptodata/1/pna/US60110_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6012_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6012_COMB.s. 5: /cgn2_6/ptodata/1/pna/US6012_COMB.s. 5: /cgn2_6/ptodata/1/pna/US6022_COMB.s. 6: /cgn2_6/ptodata/1/pna/US6022_COMB.s. 6: /cgn2_6/ptodata/1/pna/US6022_COMB.s. 6: /cgn2_6/ptodata/1/pna/US6023_COMB.s. 6: /cgn2_6/ptodata/1/pna/US6023_COMB.s. 6: /cgn2_6/ptodata/1/pna/US6023_COMB.s.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | <pre>gn2_6/ptodata/1/pna/US6026_COMB.s</pre> | 70:         |  |
| 4: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: //cgn2_6/ptodata/1/pna/US6002_COMB.s. 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s. 0: //cgn2_6/ptodata/1/pna/US6006_COMB.s. 0: //cgn2_6/ptodata/1/pna/US60010_COMB.s. 0: //cgn2_6/ptodata/1/pna/US6010_COMB.s. 2: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 3: //cgn2_6/ptodata/1/pna/US60110_COMB.s. 5: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 6: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 6: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 6: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 6: //cgn2_6/ptodata/1/pna/US60113_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60115_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60115_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60115_COMB.s. 9: //cgn2_6/ptodata/1/pna/US60115_COMB.s. 9: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 0: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 0: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 0: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 0: //cgn2_6/ptodata/1/pna/US6012_COMB.s. 0: //cgn2_6/ptodata/1/pna/US6012_COMB.s.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | gn2_6/ptodata/1/pna/US6025_COMB.s            | 69:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: /cgn2_6/ptodata/1/pna/US6001_COMB.s. 6: /cgn2_6/ptodata/1/pna/US6003_COMB.s. 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6008_COMB.s. 1: /cgn2_6/ptodata/1/pna/US6008_COMB.s. 2: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 3: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 4: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6012_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6022_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6022_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6022_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6022_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6022_COMB.s. 9: /cgn2_6/ptodata/1/pna/US6022_COMB.s.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        | <pre>gn2_6/ptodata/1/pna/US6024_COMB.s</pre> | 68:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6001_COMB.s 5: /cgn2_6/ptodata/1/pna/US6001_COMB.s 6: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US60006_COMB.s 1: /cgn2_6/ptodata/1/pna/US60010_COMB.s 1: /cgn2_6/ptodata/1/pna/US6010_COMB.s 3: /cgn2_6/ptodata/1/pna/US6011_COMB.s 5: /cgn2_6/ptodata/1/pna/US6011_COMB.s 8: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6012_COMB.s 9: /cgn2_6/ptodata/1/pna/US6012_COMB.s 9: /cgn2_6/ptodata/1/pna/US6012_COMB.s 9: /cgn2_6/ptodata/1/pna/US6013_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 | 9n2_6/ptodata/1/pna/US6023_COMB.s            | 67:         |  |
| 4: //cgn2_6/ptodata/1/pna/US6000_COMB.s<br>6: //cgn2_6/ptodata/1/pna/US6001_COMB.s<br>6: //cgn2_6/ptodata/1/pna/US6003_COMB.s<br>7: //cgn2_6/ptodata/1/pna/US6003_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6006_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6006_COMB.s<br>0: //cgn2_6/ptodata/1/pna/US60009_COMB.s<br>0: //cgn2_6/ptodata/1/pna/US60010_COMB.s<br>0: //cgn2_6/ptodata/1/pna/US6010_COMB.s<br>0: //cgn2_6/ptodata/1/pna/US60110_COMB.s<br>2: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>3: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>6: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>6: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>6: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6011_COMB.s<br>9: //cgn2_6/ptodata/1/pna/US6011_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  | gn2_6/ptodata/1/pna/US6022_COMB.s            | 66:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s 6: /cgn2_6/ptodata/1/pna/US6001_COMB.s 6: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6004_COMB.s 8: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US6006_COMB.s 1: /cgn2_6/ptodata/1/pna/US60010_COMB.s 1: /cgn2_6/ptodata/1/pna/US6010_COMB.s 3: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s 1: /cgn2_6/ptodata/1/pna/US6018_COMB.s 1: /cgn2_6/ptodata/1/pna/US6018_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         | gn2_6/ptodata/1/pna/US6021_COMB.s            | 65:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s 5: /cgn2_6/ptodata/1/pna/US6001_COMB.s 6: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US6001_COMB.s 1: /cgn2_6/ptodata/1/pna/US6010_COMB.s 1: /cgn2_6/ptodata/1/pna/US6011_COMB.s 3: /cgn2_6/ptodata/1/pna/US6011_COMB.s 5: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      | <pre>gn2_6/ptodata/1/pna/US6020_COMB.s</pre> | 64:         |  |
| 4: //cgn2_6/ptodata/1/pna/US6000_COMB.s 5: //cgn2_6/ptodata/1/pna/US6001_COMB.s 6: //cgn2_6/ptodata/1/pna/US6003_COMB.s 7: //cgn2_6/ptodata/1/pna/US6003_COMB.s 7: //cgn2_6/ptodata/1/pna/US6004_COMB.s 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s 9: //cgn2_6/ptodata/1/pna/US6006_COMB.s 0: //cgn2_6/ptodata/1/pna/US6009_COMB.s 0: //cgn2_6/ptodata/1/pna/US6009_COMB.s 0: //cgn2_6/ptodata/1/pna/US6010_COMB.s 2: //cgn2_6/ptodata/1/pna/US6010_COMB.s 3: //cgn2_6/ptodata/1/pna/US6011_COMB.s 5: //cgn2_6/ptodata/1/pna/US6011_COMB.s 6: //cgn2_6/ptodata/1/pna/US6011_COMB.s 7: //cgn2_6/ptodata/1/pna/US6011_COMB.s 9: //cgn2_6/ptodata/1/pna/US6015_COMB.s 9: //cgn2_6/ptodata/1/pna/US6016_COMB.s 9: //cgn2_6/ptodata/1/pna/US6016_COMB.s 9: //cgn2_6/ptodata/1/pna/US6016_COMB.s 9: //cgn2_6/ptodata/1/pna/US6016_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | <pre>gn2_6/ptodata/1/pna/US6019_COMB.s</pre> | 63:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6001_COMB.s 6: /cgn2_6/ptodata/1/pna/US6001_COMB.s 6: /cgn2_6/ptodata/1/pna/US6002_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6004_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US6000_COMB.s 0: /cgn2_6/ptodata/1/pna/US6000_COMB.s 0: /cgn2_6/ptodata/1/pna/US6010_COMB.s 2: /cgn2_6/ptodata/1/pna/US6011_COMB.s 2: /cgn2_6/ptodata/1/pna/US6011_COMB.s 7: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 0: /cgn2_6/ptodata/1/pna/US6011_COMB.s 0: /cgn2_6/ptodata/1/pna/US6011_COMB.s 0: /cgn2_6/ptodata/1/pna/US6011_COMB.s 0: /cgn2_6/ptodata/1/pna/US6011_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           | gn2_6/ptodata/1/pna/US6018_COMB.s            | 62:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s 5: /cgn2_6/ptodata/1/pna/US6001_COMB.s 6: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6004_COMB.s 8: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US6008_COMB.s 1: /cgn2_6/ptodata/1/pna/US6008_COMB.s 1: /cgn2_6/ptodata/1/pna/US6010_COMB.s 3: /cgn2_6/ptodata/1/pna/US6011_COMB.s 3: /cgn2_6/ptodata/1/pna/US6011_COMB.s 5: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  | qn2_6/ptodata/1/pna/US6017 COMB s            | 61:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s 5: /cgn2_6/ptodata/1/pna/US6001_COMB.s 6: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US60010_COMB.s 1: /cgn2_6/ptodata/1/pna/US6010_COMB.s 2: /cgn2_6/ptodata/1/pna/US6011_COMB.s 3: /cgn2_6/ptodata/1/pna/US6011_COMB.s 5: /cgn2_6/ptodata/1/pna/US6011_COMB.s 8: /cgn2_6/ptodata/1/pna/US6011_COMB.s 8: /cgn2_6/ptodata/1/pna/US6011_COMB.s 9: /cgn2_6/ptodata/1/pna/US6011_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               | n2_6/ptodata/1/pna/US6016_COMB.s             | 60:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s  6: /cgn2_6/ptodata/1/pna/US6001_COMB.s  6: /cgn2_6/ptodata/1/pna/US6001_COMB.s  7: /cgn2_6/ptodata/1/pna/US6003_COMB.s  7: /cgn2_6/ptodata/1/pna/US6006_COMB.s  9: /cgn2_6/ptodata/1/pna/US6006_COMB.s  9: /cgn2_6/ptodata/1/pna/US6007_COMB.s  9: /cgn2_6/ptodata/1/pna/US6007_COMB.s  9: /cgn2_6/ptodata/1/pna/US6009_COMB.s  10: /cgn2_6/ptodata/1/pna/US60010_COMB.s  2: /cgn2_6/ptodata/1/pna/US60110_COMB.s  3: /cgn2_6/ptodata/1/pna/US60110_COMB.s  5: /cgn2_6/ptodata/1/pna/US60112_COMB.s  6: /cgn2_6/ptodata/1/pna/US60113_COMB.s  6: /cgn2_6/ptodata/1/pna/US60113_COMB.s  6: /cgn2_6/ptodata/1/pna/US60113_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         | n2_6/ptodata/1/pna/US6015 COMB.s             | 59:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s 5: /cgn2_6/ptodata/1/pna/US6001_COMB.s 6: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6004_COMB.s 8: /cgn2_6/ptodata/1/pna/US6005_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US6009_COMB.s 1: /cgn2_6/ptodata/1/pna/US6009_COMB.s 2: /cgn2_6/ptodata/1/pna/US6000_COMB.s 3: /cgn2_6/ptodata/1/pna/US6010_COMB.s 3: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              | gn2 6/ptodata/1/pna/US6014 COMB s            | 58:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s 5: /cgn2_6/ptodata/1/pna/US6001_COMB.s 6: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6004_COMB.s 8: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US6008_COMB.s 1: /cgn2_6/ptodata/1/pna/US6008_COMB.s 3: /cgn2_6/ptodata/1/pna/US6008_COMB.s 3: /cgn2_6/ptodata/1/pna/US6010_COMB.s 3: /cgn2_6/ptodata/1/pna/US6011_COMB.s 6: /cgn2_6/ptodata/1/pna/US6011_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              | n2 6/ptodata/1/pna/US6013 COMB s             | 57:         |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s 5: /cgn2_6/ptodata/1/pna/US6001_COMB.s 6: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6003_COMB.s 7: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 9: /cgn2_6/ptodata/1/pna/US6006_COMB.s 0: /cgn2_6/ptodata/1/pna/US6007_COMB.s 0: /cgn2_6/ptodata/1/pna/US6009_COMB.s 0: /cgn2_6/ptodata/1/pna/US6009_COMB.s 0: /cgn2_6/ptodata/1/pna/US6009_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   | n2 6/ptodata/1/pna/NS6012 Cown s             | <br>        |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s<br>5: /cgn2_6/ptodata/1/pna/US6001_COMB.s<br>6: /cgn2_6/ptodata/1/pna/US6003_COMB.s<br>7: /cgn2_6/ptodata/1/pna/US6004_COMB.s<br>8: /cgn2_6/ptodata/1/pna/US6005_COMB.s<br>9: /cgn2_6/ptodata/1/pna/US6005_COMB.s<br>9: /cgn2_6/ptodata/1/pna/US6005_COMB.s<br>0: /cgn2_6/ptodata/1/pna/US6006_COMB.s<br>1: /cgn2_6/ptodata/1/pna/US6008_COMB.s<br>2: /cgn2_6/ptodata/1/pna/US6009_COMB.s<br>3: /cgn2_6/ptodata/1/pna/US6009_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     | n2 6/ptodata/1/pna/m66011_cown 6             | л (<br>л .  |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s<br>5: /cgn2_6/ptodata/1/pna/US6001_COMB.s<br>6: /cgn2_6/ptodata/1/pna/US6003_COMB.s<br>7: /cgn2_6/ptodata/1/pna/US6003_COMB.s<br>8: /cgn2_6/ptodata/1/pna/US6004_COMB.s<br>9: /cgn2_6/ptodata/1/pna/US6005_COMB.s<br>0: /cgn2_6/ptodata/1/pna/US6006_COMB.s<br>1: /cgn2_6/ptodata/1/pna/US6006_COMB.s<br>1: /cgn2_6/ptodata/1/pna/US6008_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         | gn2 6/htcdata/1/pna/US6009_COMB.s            | л (<br>2- ( |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s<br>5: /cgn2_6/ptodata/1/pna/US6001_COMB.s<br>6: /cgn2_6/ptodata/1/pna/US6003_COMB.s<br>7: /cgn2_6/ptodata/1/pna/US6004_COMB.s<br>8: /cgn2_6/ptodata/1/pna/US6004_COMB.s<br>9: /cgn2_6/ptodata/1/pna/US6006_COMB.s<br>0: /cgn2_6/ptodata/1/pna/US6007_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             | gnz_e/prodata/i/pna/US6008_COMB.s            | л С         |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s<br>5: /cgn2_6/ptodata/1/pna/US6001_COMB.s<br>6: /cgn2_6/ptodata/1/pna/US6003_COMB.s<br>7: /cgn2_6/ptodata/1/pna/US6003_COMB.s<br>7: /cgn2_6/ptodata/1/pna/US6004_COMB.s<br>9: /cgn2_6/ptodata/1/pna/US6006_COMB.s<br>9: /cgn2_6/ptodata/1/pna/US6006_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             | gnz_b/prodata/1/pna/US6007_COMB.s            | n ∪         |  |
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| 4: /cgn2_6/ptodata/1/pna/US6000_COMB.s<br>5: /cgn2_6/ptodata/1/pna/US6001_COMB.s<br>6: /cgn2_6/ptodata/1/pna/US6003_COMB.s<br>7: /cgn2_6/ptodata/1/pna/US6003_COMB.s<br>8: /cgn2_6/ptodata/1/pna/US6004_COMB.s                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 | gn2_b/ptodata/1/pna/US6005_COMB.s            | 2 4         |  |
| 4: /cgn2_6/ptodata/1/pna/US6000_COMB. 5: /cgn2_6/ptodata/1/pna/US6001_COMB. 6: /cgn2_6/ptodata/1/pna/US6003_COMB. 7: /cgn2_6/ptodata/1/pna/US6003_COMB.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        | gn2_6/ptodata/1/pna/US6004_COMB.s            | 8:          |  |
| <pre>gn2_6/ptodata/1/pna/US6000_COMB.<br/>gn2_6/ptodata/1/pna/US6001_COMB.<br/>gn2_6/ptodata/1/pna/US6002_COMB.</pre>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          | <pre>gn2_6/ptodata/1/pna/US6003_COMB.</pre>  | 47:         |  |
| gn2_6/ptodata/1/pna/US6000_COMB.<br>gn2_6/ptodata/1/pna/US6001_COMB.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           | gn2_6/ptodata/1/pna/US6002_COMB.             | 46:         |  |
| qn2_6/ptodata/1/pna/US6000 COMB.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               | gn2_6/ptodata/1/pna/US6001_COMB.             | 45:         |  |
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.

Score Match Length DB Query

| 21                 | 20                  | 19                 | 18                 | 17                 | Ισ                | , L               | . <u>.</u> 4     | ı<br>W             | 2                 | 11                 | c 10              |                   | . 00              | 7                 | 0                  | ı U               | 4                 | ٠ س                 | 2                 | , <u>, , , , , , , , , , , , , , , , , , </u> | No.          |
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| 88.8               | 88.8                | 88.8               | 88.8               | 88.8               | 88.8              | 88.8              | 88.8             | 88.8               | 88.8              | 88.8               | 97.0              | 97.0              | 97.0              | 97.0              | 97.0               | 100.0             | 100.0             | 100.0               | 100.0             | 100.0                                         | Match        |
| 1917               | 1917                | 1917               | 1917               | 1917               | 1866              | 1866              | 1866             | 1832               | 1608              | 1608               | 43729             | 1897              | 1897              | 1897              | 1897               | 625               | 625               | 625                 | 625               | 625                                           | Match Length |
| 25                 | 21                  | 25                 | 25                 | 17                 | 74                | 73                | 42               | 40                 | 43                | Н                  | 70                | 42                | 36                | 31                | 1                  | 42                | 36                | u                   | 31                | ᆫ                                             | BB           |
| US-09-652-128-7518 | 115-09-652-121-6191 | US-09-652-109-9370 | US-09-644-873-9077 | US-09-371-168-7188 | US-60-308-868-26  | US-60-295-262-26  | US-10-208-408-26 | US-10-108-260A-197 | US-10-264-049-111 | PCT-US01-18569-111 | US-60-261-974-9   | US-10-212-677-214 | US-09-970-966-214 | US-09-825-294-214 | PCT-US01-45395-214 | US-10-212-677-210 | US-09-970-966-210 | US-09-867-701-10876 | US-09-825-294-210 | PCT-US01-45395-210                            | ID           |
| Sequence 7518, Ap  |                     |                    |                    | 7188               | Sequence 26, Appl | Sequence 26, Appl | ~                |                    | 111,              |                    | Sequence 9, Appli | 21                | 21                | Sequence 214, App | 14                 | Sequence 210, App | Sequence 210, App | 26                  | 0                 | Sequence 210, App                             | Description  |

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RESULT 1

PCT-USO1-45395-210

Sequence 210, Application PC/TUSO145395

Sequence 210, Application PC/TUSO145395

GENERAL INFORMATION:

APPLICANT: COrixa Corporation

APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

APPLICANT: Hing, Steven P.

APPLICANT: Molesh, David Alan

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.48401PC

CURRENT APPLICATION NUMBER: PCT/USO1/45395

CURRENT FILING DATE: 2001-11-13

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 210

LENGTH: 625
                                                                                                                                                                                                               ; FEATURE:
NAME/KEY: misc_feature
; LOCATION: 607
OTHER INFORMATION: n =
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                                                                                                                                 AGTTCTCCTTGCAGAGGACTGGCGCCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
                       CAGCGACAGGCGCACACCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC
                                    CAGCGACAGGCGGCACCACACACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC
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US-09-652-121-7293
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Pred. No. 6.3e-125;
Mismatches 0;
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Sequence 23, Appl
Sequence 1915, Ap
Sequence 2415, Ap
Sequence 16, Appl
Sequence 203, App
Sequence 205, App
Sequence 211, App
Sequence 205, App
Sequence 205, App
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APPLICANT: Xbolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Filog, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CA
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 210
LENGTH: 625
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TYPE: DNA
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                        CAGCGACAGGCGGCAGCACCAGCACCTGCACGACCACCCGCCGAAACTGCTGCGAGGACAC
                                                          GCCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
                                                                                                                                                        625;
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              CAGCGACAGGCGCCAGCACCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACAC
                                                                                                          AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
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                                                                                                                                                         Conservative
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                                                                                                                                                      Score 625; DB 31;
Pred. No. 6.3e-125;
Mismatches 0;
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-ب \_ Query Match Best Local Similarity Matches 625; Conserv

TYPE: DNA

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Sequence 10876, Application US/09867701

Sequence 10876, Application US/09867701

APPLICANT: Aglate, Paul A.

APPLICANT: Jones, Robert

APPLICANT: Harlocker, Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.497

CURRENT APPLICATION NUMBER: US/09/867,701

CURRENT FILING DATE: 2001-05-29

NUMBER OF SEQ ID NOS: 10912

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 10876

LENGTH: 625

TYPE: DNA

ORGANISM: Homo sapiens

FEATURE:

NAME XEVY: TASC foature
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                                                                                                                                                                                                      ; NAME/KEY: misc_feature
; LOCATION: 607
; OTHER INFORMATION: n =
US-09-867-701-10876
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APPLICANT: XU, Jangchun

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.484C6

CCURRENT APPLICATION NUMBER: US/09/970,966

CCURRENT FILING DATE: 2001-10-02

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 210

LENGTH: 625

TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: 607
OTHER INFORMATION: n = A,T,C or G
US-09-970-966-210
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                                                                                                                                                  Query Match
Best Local Similarity
Matches 625; Conserv
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APPLICANT: Molesh, David Ali
APPLICANT: Fling, Steven P.
APPLICANT: Xu, Jiangchun
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                                                                                                                                                  Conservative
                                                                                                                                                 100.0%; Score 625; DB 36;
100.0%; Pred. No. 6.3e-125;
tive 0; Mismatches 0;
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GENERAL INFORMATION:
APPLICANT: Corixa Corporation
                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 214
LENGTH: 1897
                                                                                                                                          Query Match
Best Local Similarity
Matches 606; Conserv
                                                                                                                                                                                                                                                                                    APPLICANT: Algaté, Paul A.
APPLICANT: Eling, Steven P.
APPLICANT: Wolesh, David Alan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.48401BER: PCT/US01/45395
CURRENT APPLICATION NUMBER: PCT/US01/45395
CURRENT FILING DATE: 2001-11-13
                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Xu, Jiangchun APPLICANT: Stolk, John A.
                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
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ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                               Windows Version 4.0
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                                                                                                                                            Score 606; DB 1; L
Pred. No. 5.3e-121;
0; Mismatches 0;
                                                                                                                                                                    Length 1897;
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| ?1 CAGCGACAGGCGGCAGCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC                                                                            | 12                                              | δŌ       |
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| 1 GGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG<br>                                                                         | 6 6                                             | g<br>VQ  |
| 1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAG                                                                                |                                                 | 4d<br>4o |
| / Match 100.0%; Score 625; DB 42; Length 625; Local Similarity 100.0%; Pred. No. 6.3e-125; nes 625; Conservative 0; Mismatches 0; Indels 0 | Query M<br>Best Lo<br>Matches                   |          |
| TYPE: DNA ORGANISM: Homo sapiens FEATURE: NAME/KEY: misc_feature LOCATION: 607 OTHER INFORMATION: n = A,T,C or G IS-10-212-677-210         | TYPE ORGA FEAT NAME LOCA OTHE S-10-21           | g        |
| ER OF SEQ ID NOS: 288  ARE: FastSEQ for Windows Version 4.0  NO 210  STH: 625                                                              | NUMBE<br>SOFTW<br>SEQ ID<br>LENG                |          |
| OF OVARIAN CANCER<br>212,677                                                                                                               | TITLE<br>FILE<br>CURRE                          |          |
| ND METHODS FOR TH                                                                                                                          | APPLI<br>APPLI<br>APPLI<br>TITLE                |          |
| ;<br>(2-677-210<br>ICE 210, Application US/10212677<br>ICANT: Chenault, Ruth A.                                                            | ESULT 5<br>S-10-21<br>Sequen<br>GENERA<br>APPLI |          |
| )1 ACAGTGNAAAAAAAAAAAAAAAAA 625                                                                                                            | 60                                              | 밁        |
|                                                                                                                                            | y 601                                           | Ş        |
| AACACTGTTTCAAACCCGGGGGAGTTGGCTGTGTTAAAGAAAG                                                                                                | y 541<br>b 541                                  | å å      |
| 31 GTACCAATTTTTCAGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAG                                                                                      | b 481                                           | 밁        |
| -                                                                                                                                          | 4 8                                             | δÕ       |
|                                                                                                                                            | 42                                              | 요 :      |
|                                                                                                                                            | 4 6                                             | 5 5      |
| I TGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAG<br>                                                                         |                                                 | 5 5      |
| ) CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC                                                                             | b 30                                            | B 5      |
| 0-6                                                                                                                                        | 24 4                                            | B 5      |
| , C                                                                                                                                        | - α                                             | 2 5      |
| 11 CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGG                                                                              |                                                 | 5 5      |
|                                                                                                                                            |                                                 |          |

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; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(1897)
; OTHER INFORMATION: n = A,T,C or G
US-09-825-294-214
                                                                                                                                                                                                                                                                                                  APPLICANT: Xu, Jiangchun

APPLICANT: Stolk, John A.

APPLICANT: Algate, Paul A.

APPLICANT: Fling, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE

TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.484C5

CURRENT APPLICATION NUMBER: US/09/825,294

CURRENT FILING DATE: 2001-04-03

NUMBER OF SEQ ID NOS: 215

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 214

LENGTH: 1897
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                          GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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                                                                                                                                              Conservative
                                                                                                                                                        97.0%; Score 606; DB 31; Length 100.0%; Pred. No. 5.3e-121;
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APPLICANT: Stolk, John A.

APPLICANT: Molesh, David Alan
APPLICANT: Wilesh, David Alan
APPLICANT: Fling, Steven P.
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C6
CURRENT FILLICATION NUMBER: US/09/970,966
CURRENT FILLIC DATE: 2001-10-02
NUMBER OF SEQ ID NOS: 215
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 214
LENGTH: 1897
TYPE: DNA
ORGANISM: Homo sapiens
US-09-970-966-214
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100.0%; Pred. No. 5.3e-121;
tive 0; Mismatches 0;
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NAME/KEY: misc_feature

; LOCATION: (1)...(43729)

; OTHER INFORMATION: n = A,T,C

US-60-261-974-9
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CURRENT FILING DATE: 2001-01-17
NUMBER OF SEQ ID NOS: 237
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TYPE: DNA
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100.0%; Pred. No. 2.2e-121;
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| P<br>S         | 181<br>1451                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAACGTCTCCGAGAAGGGGAG 240                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |  |
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| Db<br>Qy       | 241<br>1511                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC 300                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |  |
| do<br>Qy       | 301<br>1571                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC 360                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |  |
| D <sub>y</sub> | 361<br>1631                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | TGGGCAGACACGAGCAGGAGGGGAGAGACAGAGAAAAAAAA                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     |  |
| g<br>94        | 421<br>1691                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |  |
| 0y<br>0y       | 481<br>1751                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | GGTACCAATTTTTCAGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAG                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |  |
| 84<br>64       | 541<br>1811                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | AACACTGTTTCAAACCCGGGGGAGTTGGCTGTGTTAAAGAAAG                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   |  |
| ρ<br>γ         | 601<br>1871                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | ACAGTG 606<br>      <br> ACAGTG 1876                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |  |
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| Дu<br>Мa       | uery Ma<br>est Loc<br>atches                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   | tch 97.0%; Score 606; DB 42; Length 1897;<br>xal Similarity 100.0%; Pred. No. 5.3e-121;<br>606; Conservative 0; Mismatches 0; Indels 0; Gaps 0;                                                                                                                                                                                                                                                                                                                                                                                                                                                               |  |
| g<br>94        | 1<br>1271                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      | AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60<br>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |  |
| P<br>Oy        | 61<br>1331                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     | GGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120<br>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |  |
| ρ <sub>ο</sub> | 121<br>1391                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | CAGCGACAGGCGCAGCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC 180<br>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |  |
| Qy             | 181                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            | CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG 240                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |  |

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TITLE OF INVENTION: NUCLEAC ACIDS, PROTEINS and FILE REFERENCE: PA133CT CURRENT APPLICATION NUMBER: PCT/USO1/18569 CURRENT FILING DATE: 2001-06-07 PRIOR APPLICATION NUMBER: 60/209,467 PRIOR FILING DATE: 2000-06-07 NUMBER OF SEQ ID NOS: 4360 SOFTWARE: PATENTIN Ver. 2.0 SEQ ID NO 111 SEQ ID NO 111 LENGTH: 1608
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GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc.
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LOCATION: (22)
OTHER INFORMATION: n equals a,t,g,
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LOCATION: (87)
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LOCATION: (32)
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GGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 1044
                                                       AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 984
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                                                                                                                     Conservative
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Pred. No. 4.6e-110;
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RESULT 12
US-10-264-049-111
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GENERAL INFORMATION:
APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
                                                                                                                                                                                                                                     SEQ ID NO 111
LENGTH: 1608
                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/209,467
PRIOR FILING DATE: 2000-06-07
NUMBER OF SEQ ID NOS: 4360
SOFTWARE: Patentin Ver. 3.1
                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/264,049 CURRENT FILING DATE: 2002-10-04
                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: PCT/US01/18569 PRIOR FILING DATE: 2001-06-07
                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: PA133P1
                                                                                                                                                                                                                         TYPE: DNA
NAME/KEY: misc_feature
LOCATION: (87)..(87)
OTHER INFORMATION: n equals a,t,g, or
                                                                                                   NAME/KEY: misc_feature
LOCATION: (22)..(22)
OTHER INFORMATION: n equals
                                                    OTHER INFORMATION: n equals
                                                                  NAME/KEY: misc_feature LOCATION: (32)..(32)
                                                                                                                                                                      FEATURE:
NAME/KEY: misc_feature
LOCATION: (18)..(18)
                                                                                                                                                                                                          ORGANISM: Homo sapiens
                                                                                           FEATURE
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                                                                                                                                                          OTHER INFORMATION: n equals
                                            FEATURE
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US-10-264-049-111

Query Match Best Local

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US-10-108-260A-197

Sequence 197, Application US/10108260A

GENERAL INFORMATION:
APPLICANT: HELIX RESEARCH INSTITUTE
APPLICANT: HELIX RESEARCH INSTITUTE
TITLE OF INVENTION: Novel full length cDNA
FILE REFERENCE: H1-A0106
CURRENT APPLICATION NUMBER: US/10/108,260A
CURRENT FILING DATE: 2002-03-27
NUMBER OF SEQ ID NOS: 5458
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 197
LENGTH: 1032
TYPE: DNA
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Pred. No. 4.6e-110;
                                    Score 555; DB 40;
Pred. No. 4.4e-110;
0; Mismatches 1;
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; LENGTH: 1866
; TYPE: DNA
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GENERAL INFORMATION:
APPLICANT: Schebye, Xiao Min
TITLE OF INVENTION: GENES REGULATED BY PEROXISOME PROLIFERATOR-ACTIVATED RECEPTOR
FILE REFERENCE: PA-0048-1 US
FILE REFERENCE: PA-0048-1 US
                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                             CURRENT FILING DATE: 2002-07-29
PRIOR APPLICATION NUMBER: 60/308,868
PRIOR FILING DATE: 2001-07-30
NUMBER OF SEQ ID NOS: 55
                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PERL Program
                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                              OTHER INFORMATION: Incyte ID
                                                                                                                                                                                                                  NAME/KEY: misc_feature
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                                                                                                                                                                                                NO:
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                                                                                                          Score 555; DB 42;
Pred. No. 4.4e-110;
                                                                                        Mismatches
                                                                                                                                                                                                  227484.13
                                                                                                                                   Length 1866;
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AGTTCTCCTTGCAGAGGACTGGCCCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG

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; ORGANISM: Homo sapiens US-10-108-260A-197

Query Match
Best Local Similarity

Matches

605;

Conservative

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Sequence 26, Application US/60295262
(GENERAL INFORMATION:
APPLICANT: Schebye, Xiao Min
TITLE OF INVENTION: GENES REGULATED BY PEROX
TITLE OF INVENTION: GAMMA AGONIST
FILE REFERENCE: PA-0048 P
CURRENT APPLICATION NUMBER: US/60/295,262
CURRENT FILING DATE: 2001-06-01
NUMBER OF SEQ ID NOS: 51
SOFTMARE: PERL PROGram
SEQ ID NO 26
LENGTH: 1866
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                                                                                                                                      ; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID
US-60-295-262-26
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US-60-295-262-26
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ORGANISM: Homo sapiens
FEATURE:
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                                              1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60
GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                  AGTTCTCCTTGCAGAGGACTGGCGCCCGGACGCGAAGAGCAACGGCCCTGCACAAAGCG 1200
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                                                                                        605;
                                                                                                  Similarity
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                                                                                      Conservative
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                                                                                               Score 555; DB 73;
Pred. No. 4.4e-110;
                                                                                    Mismatches
                                                                                                                                                227484.13
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                                                                                                          Length 1866;
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US-60-308-868-26
; Sequence 26, Application US/60308868
; Sequence 26, Application US/60308868
; Sequence 26, Application US/60308868
; Sequence 26, Application US/60308868
; TITLE OF INFORMATION: GENES REGULATED BY PEROXISOME PROLIFERATOR-ACTIVATED RECEPTOR
; FILE REFERENCE: PA-0048-1 P
; CURRENT APPLICATION NUMBER: US/60/308,868
; CURRENT FILING DATE: 2001-07-30
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SOFTWARE: PERL Program
SEQ ID NO 26
LENGTH: 1866
TYPE: NO.
                                                                                                                                                     Matches 605;
                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                   NAME/KEY: misc_feature
OTHER INFORMATION: Incyte
                                                                                                                                                                                                                                                 ORGANISM: HOMO FEATURE:
                                                                                                                                                                                                                                                                             TYPE: DNA
                                                                                            1141 AGTTCTCCTTGCAGAGGACTGGCGCGCGGGACGCGAAGAGCAACGGGGCGCTGCACAAAGCG 1200
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                                                                                                         CAGCGACAGGCGGCAGCACCAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC
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Conservative
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                                                                                                                                                               99.8%;
                                                                                                                                                                                                                      ID No:
                                                                                                                                                   0;
                                                                                                                                                 Score 555; DB 74;
Pred. No. 4.4e-110;
0; Mismatches 1;
                                                                                                                                                                           Length 1866;
                                                                                                                                                   Indels
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; SEQ ID NO 9077
; LENGTH: 1917
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-644-873-9077
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                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 605; Conservat
                                                                                                                                                                                                                                                                                                                            APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: Holtzman, Douglas A.
TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND
TITLE OF INVENTION: THEREFOR
TILE REFERENCE: 1600.1169-001
CURRENT APPLICATION NUMBER: US/09/644,873
CURRENT FILING DATE: 2000-08-28
PRIOR APPLICATION NUMBER: 60/151,064
PRIOR APPLICATION NUMBER: 60/151,064
PRIOR FILING DATE: 1999-08-27
NUMBER OF SEQ ID NOS: 11286
TOURNELL TO THE SEQ ID NOS: 11286
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                                                                                                                                                                   1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60
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                                                                                                                GGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
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CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGGAG
                                               CAGCGACAGGCGGCAGCACCACCTGCACGAACACCCGGCCGAAACTGCTGCGAGGACAC
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                                                                                                                                                                                                      Score 555; DB 25;
Pred. No. 4.4e-110;
0; Mismatches 1;
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| CAGCGACAGGCGGCAGCACCAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC 180 | GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGGGCTTCTCGTGGTTGGCGTGCTG 120<br> | AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60<br> | / Match 88.8%; Score 555; DB 17; Length 1917;<br>Local Similarity 99.8%; Pred. No. 4.4e-110;<br>nes 605; Conservative 0; Mismatches 1; Indels 0; Gaps | ESULT 17 S-09-371-168-7188 Sequence 7188, Application US/09371168 GENERAL INFORMATION: APPLICANT: Glucksmann, M. Alexandra TITLE OF INVENTION: NUCLEIC ACID MOLECULES DERIVED FROM A TITLE OF INVENTION: HUMAN BRAIN LIBRARY FILE REFERENCE: MLN98-37pA CURRENT APPLICATION NUMBER: US/09/371,168 CURRENT FILING DATE: 1999-08-10 EARLIER APPLICATION NUMBER: 60/095,907 EARLIER APPLICATION NUMBER: 60/103,145 EARLIER FILING DATE: 1998-08-10 EARLIER FILING DATE: 1998-10-05 NUMBER OF SEQ ID NOS: 8285 SOFTWARE: FASTSEQ for Windows Version 3.0 SEQ ID NO 7188 LENGTH: 1917 TYPE: DNA ORGANISM: Homo Sapiens                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              | ACAGTG 1746 | ACAGTG 606 | AACACTGTTTCAAACCCGGGGGAGTTGGCTGTGTTAAAGAAAG | AACACTGTTTCAAACCCCGGGGGAGTTGGCTGTGTTAAAGAAAAGACCATTAAATGCTTTTAG 600 | 6 4  |     | TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480 | TGGGCAGACACGAGGAGGAGAGAGAGAAAAGAAAAACACAGCATGAGAACACAG 1560 | TGGGCAGACACGAGCAGGAGAGAGAGAGAGAAAAGAAAA | CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC 360 | C 1  | GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC 300 | CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAACGTCTCCGAGAAGGGGAG 1380 | CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGGAG 240 | CAGCGACAGGCGGCAGCACCAGCACCAGCAACACCCGGCGAAACTGCTGCGAGGACAC 1320 |

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Sequence 9370, Application US/09652109

GENERAL INFORMATION:
APPLICANT: MCGATTHY, Sean A.
TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
TITLE OF INVENTION: THEREFOR
FILE REFERENCE: 1600.1180-001
CURRENT APPLICATION NUMBER: US/09/652,109
CURRENT FILING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/151,128
PRIOR FILING DATE: 1999-08-30
NUMBER OF SEQ ID NOS: 10105
SOFTMARE: FastSEQ for Windows Version 4.0
LENGTH: 1917
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              GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC 300
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Pred. No. 4.4e-110;
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CURRENT APPLICATION NUMBER: US/09/652,121
CURRENT FILING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/151,129
PRIOR FILING DATE: 1999-08-30
NUMBER OF SEQ ID NOS: 7615
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 6191
LENGTH: 1917
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Sequence 6191, Application US/09652121
GENERAL INFORMATION:
APPLICANT: Distefano, Peter
TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
TITLE OF INVENTION: THEREFOR
FILE REFERENCE: 1600.1188-001
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| Db 1646 TGGGCAGACACGAGCAGGAGGAGAGAGAAAAAGAAAAACACAGCATGAGAACACAG 1705  Qy 421 TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480 | 361 TGGGCAGACAGGAGCAGGAGGAGAGAGAGAAAAGAAAAACAGAGCATGAGAACACAG | Qy 301 CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC 360<br> | QY 241 GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC 300 | QY 181 CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCCGAGAAGGGGAG 240 | OY 121 CAGCGACAGGCGCAGCACAGCACCTGCACGACACCCGCCGAAACTGCTGCGAGGACAC 180 | Qy 61 GGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120 | QY 1 AGTICTCCTTGCAGAGGACTGGCGCCCGGGACGCGAAGAGCCAACGGGCGCTGCACAAAGCG 60 | Query Match 88.8%; Score 555; DB 25; Length 1917; Best Local Similarity 99.8%; Pred. No. 4.4e-110; Matches 605; Conservative 0; Mismatches 1; Indels 0; Gaps 0; | OY 421 TAAATGAATAAAAAAAAAAAAAAAAAAAAAAAAAAAAA |

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| 361 TGGGCAGACACGAGCAGGAGGGAGGACAGAGAAAAGAAA | 301 CATGATCCTCCGAATCTGGTTGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC 36 | 241 GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC 30 | 181 CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG 24<br> | 121 CAGCGACAGGCGGCAGCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC 18 | 61 GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGCGCTTCTCGTGGTTGGCGTGCTG 12<br> | 1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60<br> | Query Match 88.8%; Score 555; DB 25; Length 1917; Best Local Similarity 99.8%; Pred. No. 4.4e-110; Matches 605; Conservative 0; Mismatches 1; Indels 0; Gaps | Sequence 7518, Application US/09652128  GENERAL INFORMATION: APPLICANT: Shylan, Andrew W. APPLICANT: Shylan, Douglas A. TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES TITLE APPLICATION NUMBER: US/09/652,128 CURRENT FILING DATE: 2000-08-30 PRIOR FILING DATE: 2000-08-30 PRIOR FILING DATE: 1999-08-30 PRIOR FILING DATE: 1999-08-30 NUMBER OF SEQ ID NOS: 10265 SOSTWARE: FASTSEQ for Windows Version 4.0 SEQ ID NO 7518 LENGTH: 1917 TYPE: DNA ORGANISM: Homo Sapiens | SULT 21 | 1886 ACAGTG 1891 | 541 AACACTGT | 481 GGTACCAATTTTTCAGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAGAGAGAATTT 54<br> | 421 TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 48 | 361 TGGGCAGACACGAGCAGGAGGAGGAGAGAGAAAAGAAAA |
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SEQ ID NO 8473
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TITLE OF INVENTION: MOVEL NUCLEIC ACID MOTEL NUCLEIC ACID MOTEL OF INVENTION: THEREFOR
FILE REFERENCE: 1600.1193-001
CURRENT APPLICATION NUMBER: 05/09/652,914
CURRENT FILING DATE: 2000-08-31
PRIOR APPLICATION NUMBER: 60/152,112
PRIOR FILING DATE: 1999-08-31
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           CATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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                                      TAAATAAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 1765
                                                    TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480
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                                                                                            GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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SEQ ID NO 2477
LENGTH: 1917
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PRIOR FILING DATE: 1999-08-30
NUMBER OF SEQ ID NOS: 3855
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                                                       TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480
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             CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
TGGGCAGACACGAGCAGGAGAGAGAGAAAAAGAAAAACACAGCATGAGAACACAG
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Pred No. 4.4e-110;
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PRIOR FILING DATE: 1998-09-15
PRIOR APPLICATION NUMBER: 60/106,443
PRIOR FILING DATE: 1998-10-30
PRIOR APPLICATION NUMBER: 60/107,257
PRIOR FILING DATE: 1998-11-05
PRIOR APPLICATION NUMBER: 60/126,906
PRIOR APPLICATION NUMBER: 60/126,906
PRIOR FILING DATE: 1999-03-30
PRIOR FILING DATE: 1999-03-30
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; SEQ ID NO 4454
; LENGTH: 1918
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Best Local Similarity
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APPLICANT: Holtzman, Douglas A.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES DERIVED
TITLE OF INVENTION: HUMAN FETAL SKIN LIBRARY
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CURRENT FILING DATE: 1999-09-15
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PRIOR FILING DATE: 1999-04-30
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                                                                                                                              CATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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Pred. No. 4.4e-110;
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GENERAL INFORMATION:

APPLICANT: GLUCKSMENT, M. Alexandra
TITLE OF INVENTION: HUMAN BRAIN LIBRARY
FILE OF INVENTION: HUMAN BRAIN LIBRARY
FILE REFERENCE: 1600.1037-005
CURRENT APPLICATION UMBER: US/09/801,833
CURRENT FILING DATE: 2001-03-13
PRIOR APPLICATION NUMBER: 09/371,168
PRIOR FILING DATE: 1999-08-10
PRIOR FILING DATE: 1998-08-09
PRIOR APPLICATION NUMBER: 60/095,907
PRIOR APPLICATION NUMBER: 60/095,907
PRIOR APPLICATION NUMBER: 60/103,145
PRIOR FILING DATE: 1998-10-05
PRIOR FILING DATE: 1998-10-05
NUMBER OF SEQ ID NOS: 8285
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Best Local
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                                                                                                                                                                                                                                                                 GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTTGGCCGCAGC
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 TGGGCAGACACGAGCAGGAGGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAG 420
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                                                                     TAAATGAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480
                                                                                                         TGGGCAGACACGAGCAGGAGGAGAGAGAAAAAGAAAAACACAGCATGAGAACACAG
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Pred. No. 4.4e-110;
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US-09-808-383-4454
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PRIOR FILING DATE: 1999-09-15
PRIOR PRIOR PLICATION NUMBER: US 60/100,465
PRIOR FILING DATE: 1998-09-15
PRIOR APPLICATION NUMBER: US 60/106,443
PRIOR FILING DATE: 1998-10-30
PRIOR APPLICATION NUMBER: US 60/107,257
PRIOR FILING DATE: 1998-11-15
PRIOR FILING DATE: 1998-11-15
PRIOR APPLICATION NUMBER: US 60/126,906
PRIOR APPLICATION NUMBER: US 60/132,099
PRIOR APPLICATION NUMBER: US 60/132,099
PRIOR FILING DATE: 1999-04-30
                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 5775
SOFTWARE: FASTSEQ for Windows Version 4.0
LENGTH: 1918
TYPF: Pair
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APPLICANT: Gearing, David P.
APPLICANT: Holtzman, Douglas A.
                                                                                                                                                                                                                                                                                                Matches
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Best Local :
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CURRENT APPLICATION NUMBER: US/09/808,383
CURRENT FILING DATE: 2001-03-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Holtzman, Douglas A.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES DERIVED
TITLE OF INVENTION: HUMAN FETAL SKIN LIBRARY
                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
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                                                                                                                                                                                                                                                                                                            Local Similarity
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GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
                                                             GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTTGGCCGCAGC
                                                                                                CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
                                                                                                             CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
                                                                                                                                               CAGCGACAGGCGGCAGCACCAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC
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                                                                                                                                                                                              GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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Pred. No. 4.4e-110;
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; TYPE: DNA; Homo sapiens US-09-652-121-7293
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CURRENT APPLICATION NUMBER: US/09/652,121
CURRENT FILING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/151,129
PRIOR FILING DATE: 199-08-30
NUMBER OF SEQ ID NOS: 7615
NUMBER OF SEQ ID NOS: 7615
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Best Local
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              TGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAAAACACAGCATGAGAACACAG
                                                         CATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
                                                                                                                    GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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TGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAAGAAAACACAGCATGAGAACACAG
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Pred. No. 4.4e-110;
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SEQ ID NO 3346
LENGTH: 1925
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 3346, Application US/09652917 GENERAL INFORMATION:
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Best Local Similarity
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CURRENT FILING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/151,422
PRIOR FILING DATE: 1999-08-30
NUMBER OF SEQ ID NOS: 3855
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                    CATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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                                                                                                              CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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RESULT 29
US-09-652-128-9375/c
US-09-652-128-9375/c
Sequence 9375, Application US/09652128
GENERAL INFORMATION:
APPLICANT: Shyjan, Andrew W.
APPLICANT: Holtzman, Douglas A.
TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
TITLE OF INVENTION: THEREFOR
FILE REFERENCE: 1600-1171-001
CURRENT APPLICATION NUMBER: US/09/652,128
CURRENT FILING DATE: 2000-08-30
PRIOR APPLICATION NUMBER: 60/151,133
PRIOR FILING DATE: 1999-08-30
ONUMBER OF SEQ ID NOS: 10265
SOCTWARE: FRATSEQ for Windows Version 4.0
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LENGTH: 1925
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Best Local
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TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
                                                                                     GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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Pred. No. 4.4e-110;
0; Mismatches 1;
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TAAATAAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT

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APPLICANT: Curtis, Rory A.J.

APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: Silos-Santiago, Inmaculada
APPLICANT: Distefano, Peter
ITILE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
ITILE OF INVENTION: THEREFOR
FILLE REFERENCE: 1600.2020-001
CURRENT FILING DATE: 2000-10-30
PRIOR APPLICATION NUMBER: 60/162,359
PRIOR APPLICATION NUMBER: 60/162,359
PRIOR FILING DATE: 1999-10-29
PRIOR FILING DATE: 1999-10-29
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11085
LENGTH: 1925
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; ORGANISM: Homo sapiens
US-09-699-997-11085
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                      CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
                                                                          TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
                                                                                                                  GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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        TAAATAAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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APPLICANT: Shyjan, Andrew W.

APPLICANT: Stbdal, Hilde

TITLE OF INVENTION: NOYEL NUCLEIC ACID MOLECULES AND USES

TITLE OF INVENTION: THEREFOR

FILE REFERENCE: 1600.2036-001

CURRENT APPLICATION NUMBER: US/09/710,281

CURRENT FILING DATE: 2000-11-10

PRIOR APPLICATION NUMBER: 60/164,254

PRIOR ETLING DATE: 1999-11-09

NUMBER OF SEQ ID NOS: 5803

SOFTWARE: FRSTSEQ for Windows Version 4.0
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-710-281-4458
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US-09-710-281-4458/c
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SOFTWARE: Fast
; SEQ ID NO 4458
FENGTH: 1925
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Best Local
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                          GGTACCAATTTITCAGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAGAGAGAATTT
                                                                                   CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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Pred. No. 4.4e-110;
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Length 1925;
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PRIOR FILING DATE: 2000-01-20
NUMBER OF SEQ ID NOS: 156
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 38
LENGTH: 1953
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; LOCATION: (1362)
; OTHER INFORMATION: n equals a,t,g, or
PCT-US00-09066-38
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: FILE REFERENCE: PS5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/128,694
PRIOR FILING DATE: 1999-04-09
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
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                                                                                                                                                        CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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                                                        TGGGCAGACACGAGCAGGAGGGAGACAGAGAAAAAAGAAAAACACAGCATGAGAACACAG
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Pred. No. 4.3e-110;
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US-09-726-805-1608/c
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SEQ ID NO 1608
LENGTH: 1925
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES TITLE OF INVENTION: THEREFOR FILE REFERENCE: 1600 2017-001 CURRENT APPLICATION NUMBER: US/09/726,805 CURRENT FILING DATE: 2000-11-30 PRIOR APPLICATION NUMBER: 60/168,140 PRIOR APPLICATION NUMBER: 60/168,140 PRIOR FILING DATE: 1999-11-30 NUMBER OF SEQ ID NOS: 2158
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                                       TAAATAAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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Pred. No. 4.4e-110;
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OY 601 ACACTG 606

Db 84 ACACTG 79

RESULT 35

Sequence 1914, Application US/09950083

APPLICANT; ROSEN, et. al

TITLE OF INVENTION: HUMBER: US/09/950,083

GENERAL INFORMATION: HUMBER: US/09/950,083

GENERAL INFORMATION: HUMBER: US/09/950,083

CURRENT FILING DATE: 2001-09-12

PRIOR APPLICATION NUMBER: US/09/950,083

CURRENT FILING DATE: 2001-03-27

PRIOR APPLICATION NUMBER: PCT/US00/06043

PRIOR APPLICATION NUMBER: PCT/US00/06043

PRIOR APPLICATION NUMBER: PCT/US00/06043

PRIOR APPLICATION NUMBER: PCT/US00/06043

PRIOR APPLICATION NUMBER: PCT/US00/06043

PRIOR APPLICATION NUMBER: PCT/US00/06044

PRIOR APPLICATION NUMBER: PCT/US00/06044

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PRIOR APPLICATION NUMBER: PCT/US00/06045

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PRIOR APPLICATION NUMBER: PCT/US00/06046

PRIOR APPLICATION NUMBER: PCT/US00/06046

PRIOR APPLICATION NUMBER: PCT/US00/06082

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PRIOR APPLICATION NUMBER: PCT/US00/06082

PRIOR APPLICATION NUMBER: PCT/US00/06082

PRIOR APPLICATION NUMBER: PCT/US00/06082

PRIOR APPLICATION NUMBER: PCT/US00/0682

PRIOR APPLICATION NUMBER: P
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US-10-105-299-2414

LOCATION: (1362)..(1362) OTHER INFORMATION: n equals a,t,g,

or

FEATURE: misc\_feature TYPE: DNA ORGANISM: Homo sapiens

Query Match Best Local Matches

Similarity

99.88;

Conservative

0;

Score 555; DB 40; Pred. No. 4.3e-110; 0; Mismatches 1;

Length 1953; Indels

0,:

Gaps

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625

NUMBER OF SEQ ID NOS: 15197
Prior Application removed SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2414
LENGTH: 1953

See

File Wrapper or

CURRENT APPLICATION NUMBER: US/10/105,299
CURRENT FILING DATE: 2002-03-26 TITLE OF INVENTION: Human Secreted Proteins FILE REFERENCE: PS950

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AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG

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PRIOR PRIOR PRIOR PRIOR PRIOR APPLICATION NUMBER: US 60/125,359 FILING DATE: 1999-03-19 FILING DATE: 1999-12-03 APPLICATION NUMBER: US 60/126,051 FILING DATE: 1999-03-23 FILING DATE: 1999-12-03 APPLICATION NUMBER: US APPLICATION NUMBER: US 60/168,666 NUMBER: US 60/169,906 60/168,664

PRIOR PRIOR FILING DATE: 1999-03-12 APPLICATION NUMBER: US 60/138,597 FILING DATE: 1999-06-11 FILING DATE: 1999-12-03
APPLICATION NUMBER: US 60/124,142

PRIOR PRIOR PRIOR OR APPLICATION NUMBER: US 60/168,662
OR FILING DATE: 1999-12-03
OR APPLICATION NUMBER: US 60/124,144
OR FILING DATE: 1999-03-12
OR APPLICATION NUMBER: US 60/138,574
OR FILING DATE: 1999-06-11
OR APPLICATION NUMBER: US 60/168,667

PRIOR PRIOR PRIOR PRIOR PRIOR FILING DATE: 1999-06-11

PRIOR PRIOR PRIOR PRIOR PRIOR

PRIOR PRIOR PRIOR OR FILING DATE: 1999-12-03
OR APPLICATION NUMBER: US 60/124,095
OR APPLICATION NUMBER: US 60/138,598
OR FILING DATE: 1999-05-11
OR APPLICATION NUMBER: US 60/168,665
OR FILING DATE: 1999-12-03
OR FILING DATE: 1999-12-03
OR FILING DATE: 1999-03-19
OR APPLICATION NUMBER: US 60/125,360
OR APPLICATION NUMBER: US 60/125,360
OR APPLICATION NUMBER: US 60/138,626

PRIOR PRIOR PRIOR FILING DATE: 1999-U3-12
APPLICATION NUMBER: US 60/168,663
APPLICATION 10MBER: US 60/168,663

PRIOR PRIOR PRIOR APPLICATION NUMBER: US 60/168,622 FILING DATE: 1999-12-03 APPLICATION NUMBER: US 60/124,143

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US-10-105-299-2414/c

Sequence 2414, Application US/10105299 GENERAL INFORMATION: APPLICANT: Rosen, et. al

PRIOR PRIOR PRIOR PRIOR PRIOR FILING DATE: FILING DATE: 1999-03-12 FILING DATE: 1999-03-12 APPLICATION NUMBER: US 60/124,096 APPLICATION NUMBER: US 60/168,661 FILING DATE: 1999-12-03

PRIOR PRIOR PRIOR PRIOR FILING DATE: 1999-12-03
APPLICATION NUMBER: US 60/124,099 FILING DATE: 1999-03-12
APPLICATION NUMBER: US 60/168,654 PRIOR PRIOR PRIOR PRIOR PRIOR

FILING DATE: 1999-03-12
APPLICATION NUMBER: US 60/167,061
FILING DATE: 1999-11-23

APPLICATION NUMBER: US 60/124,146

2001-04-27 2000-09-26

PCT/US01/13318

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APPLICATION NUMBER: US 60/124,093 FILING DATE: 1999-03-12 APPLICATION NUMBER: US 60/166,989 FILING DATE: 1999-11-23

APPLICATION NUMBER: US 60/124,145

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FILING DATE:

APPLICATION NUMBER:

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Length 1953;

FILING DATE: 2000-09-26
APPLICATION NUMBER: PCT/US00/26323
FILING DATE: 2000-09-26

APPLICATION NUMBER: PCT/US00/26337

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PCT-US00-14973-23/c Sequence 23, Application PC/TUS0014973 GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                         Matches
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                                              GGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
CAGCGACAGGCGCAGCACCAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC
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                              GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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Pred. No. 4.3e-110;
0; Mismatches 1;
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                   PRIOR PELICATION NUMBER: 60/278,650
PRIOR PELICATION NUMBER: 09/278,650
PRIOR PELICATION NUMBER: 09/833,245
PRIOR PELICATION NUMBER: 09/833,245
PRIOR PELICATION NUMBER: PCT/USO1/11988
PRIOR PELICATION NUMBER: PCT/USO1/6043
PRIOR APPLICATION NUMBER: PCT/USO1/6043
PRIOR PELICATION NUMBER: PCT/USO1/6043
PRIOR PELICATION NUMBER: PCT/USO1/6043
PRIOR PELICATION NUMBER: PCT/USO1/6058
PRIOR PELICATION NUMBER: PCT/USO1/6044
PRIOR PELICATION NUMBER: PCT/USO1/6044
PRIOR PELICATION NUMBER: PCT/USO1/6044
PRIOR PELICATION NUMBER: PCT/USO1/6044
PRIOR PELICATION NUMBER: PCT/USO1/6042
PRIOR APPLICATION NUMBER: PCT/USO1/6042
PRIOR APPLICATION NUMBER: PCT/USO1/6044
PRIOR PELICATION NUMBER: PCT/USO1/6014
PRIOR PELICATION NUMBER: PCT/USO1/6014
PRIOR PELICATION NUMBER: PCT/USO1/6014
PRIOR PELICATION NUMBER: PCT/USO1/6014
PRIOR PELICATION NUMBER: PCT/USO1/6049
PRIOR APPLICATION NUMBER: PCT/USO1/6049
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US-09-950-083-1915/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/950,083
CURRENT FILING DATE: 2001-09-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Human Secreted Proteins FILE REFERENCE: PS805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Rosen,
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| NUMBER: US: 1999-12-03  | 99-06-   | 99-03-     | 9-12<br>ER:     | MBER: US   | NUMBER: US | NUMBER: US | NUMBER: US<br>1999-12-0 | : 1999-03-1 | 9-12-    | នៈ         | NUMBER: US | NUMBER: US | NUMBER: US | : 1999-12-0 | . 1999-03-1     | NUMBER: US | NUMBER: US | NUMBER: US      | : 1999-11- | : 1999-03 | : 2001-04 | NUMBER:      | NUMBER:      | NUMBER:           | : 2000-09 | : 2000-09      | : 2000-09 | NUMBER:     | NUMBER:      | NUMBER:           | 0-06      | 2000-                                   | 2000-06    | NUMBER:      | NUMBER: P    | 0-06-0                     | 0-06-0     | 00-06-0        | BER: PC      | NUMBER: PC   | : 2000-04-06 | 2000-04-0 | NUMBER: PC:  |
| 60/168,662              | 60/138,6 |            | 8<br>60/125,360 | 60/168,665 | 60/138,598 | 60/124,095 | 60/168,663<br>3         | 2           | 60/10/14 | 60/168,622 | 60/124,096 | 60/168,661 | 2          | ) Hoo , oo  | 2<br>607168.654 | 60/124,145 | 60/166,989 | 60/124,093<br>2 | 60/16/,06  |           | 50/124.14 | r/us01/13318 | r/us00/26337 | r/us00/26323<br>s | 5         | ALCOUNT (1963) | /11500/   | r/us00/2637 | r/us00/14964 | r/us00/14973<br>1 | 1         | ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,, | L<br>      | r/US00/14933 | r/us00/14934 | r/US00/15135<br>l          | /0000/1400 | /IISO0 /1 / 06 | r/US00/14926 | r/us00/15136 | /0800/090    |           | r/us00/09071 |

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OR APPLICATION NUMBER: US 60/168,667
OR FILING DATE: 1999-12-03
OR APPLICATION NUMBER: US 60/124,142
OR FILING DATE: 1999-03-12
OR APPLICATION NUMBER: US 60/138,597
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; NAME/KEY: misc_feature ; LOCATION: (1362)..(1362) ; OTHER INFORMATION: n equals a,t,g, US-10-105-299-2415
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NUMBER OF SEQ ID NOS: 15197
Prior Application removed - See I
SOFTWARE: Patentin Ver. 2.0
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                                        GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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RESULT 40 PCT-US02-29964-16

Sequence 16, Application PC/TUS0229964
GENERAL INFORMATION:
APPLICANT: Tang, Y. Tom
APPLICANT: Ren, Feiyan

| g<br>Qy                   | Qy<br>Db                            | g<br>V                    | Qy<br>db                                                    | DP<br>QA                         |                      | Ğ                          |               |         |          |                                                    |       |        |          |                |            |        |            |       |                |            |      |            |                          |              |      |        |                 |             |
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| 241<br>1589               | 181<br>1529                         | 121<br>1469               | 61<br>1409                                                  | 134                              | ry M<br>t Lo<br>ches | NAME/I<br>LOCAT<br>I-US02- | YPE:<br>RGANI | OI W.E. | ma i     | IOR                                                |       |        |          | I GR           |            |        | R I G      | ROR   | ROI            | RREI       |      | PLIC       | PLIC<br>PLIC             | PLIC<br>PLIC | PITC | 110    | PLIC            | APPLICANT:  |
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| 8=8                       | GGT                                 | AGC.                      | 166                                                         | AGG<br>AGG                       | at 198               | 3)                         | ns            | Ve      | 99<br>99 | BER<br>00-                                         | BER   | BER    | BER<br>- | BER<br>01-     | BER<br>00- | BER    | BER<br>00- | BER   | BER            | 20<br>20   | IP 1 | Rado       | ent                      | .₽Ţ          | Tom  | Malabi | ω,              | •           |
| 668<br>                   | TGAT                                | ACAG<br>    <br> ACAG     | AGTG<br>    <br> AGTG                                       | BAGGACTGG<br>BAGGACTGG           | 9.8<br>9.8           |                            |               | rsi     | 2 on     | PLICATION NUMBER: US 09/5<br>LING DATE: 2000-02-28 | 3 ° 6 | 04 - I | 02-(     | 01 -:<br>01 -: | 21         | 13 P   | 04 - :     | 01 U. | . U            | 02-<br>02- | PCT  | L O G      | υ,                       | le W         | u    | ika    | ٠ <u>۽</u>      | <b>.</b> .  |
| AAGT/                     | rgac<br>       <br>  GAC            | GCAC                      | 3000<br>                                                    | 9006<br>                         | ** **<br>O           |                            |               | on 6    | dat      | 28 09                                              | 2년:   | 3709   | 300      | T/US           | 5 6        | S II I | 500        | 3 09  | 19<br>19<br>19 | <br>P.C.   | otic | ICT6       | ana                      |              |      |        |                 |             |
| AGGA<br>         <br>AGGA | CGA                                 | CIG                       | ATG                                                         | 9009                             | Sco<br>Pre           |                            |               | .0      | tar      | 9/51                                               | /US01 | /56    | )/49     | JS01           | )/49       | JS00   | /55        | /48   | )/32           | sn/c       | les  | Ċ          | _                        |              |      |        |                 |             |
| ACCT                      | GCT GCT                             | SAC   CAC                 | TAC                                                         | GGAC                             | re<br>d.<br>Mis      |                            |               |         | ето      | 5,1                                                | /03   | 0,8    | 6,9      | /02            | 1,4        | /35    | 2,3        | 8,7   | 3,7            | 02/        |      | Aci        |                          |              |      |        |                 |             |
| CGTC                      | GAG<br>     <br>GAG                 | GAAC<br>       <br>  GAAC | 1000<br>1000<br>1000<br>1000<br>1000<br>1000<br>1000<br>100 | 8 <del>-</del> 8                 | No.                  |                            |               |         | ved      | 26                                                 | 800   | 75     | 14       | 623            | 04         | 017    | 17         | 25    | 39             | 299        |      | S.         |                          |              |      |        |                 |             |
| CCAG                      | GTAG<br>    <br>GTAG                | 8-8                       | AG – AG                                                     | GAAG.                            | , 4.<br>che          |                            |               |         |          |                                                    |       |        |          |                |            |        |            |       |                | 64         |      | and        |                          |              |      |        |                 |             |
| G - G                     | GAA<br>I - A                        | 2999<br>                  | 6060<br>                                                    | GAGC<br>1111<br>GAGC             | DB 1<br>3e-1<br>s    |                            |               |         | See      |                                                    |       |        |          |                |            |        |            |       |                |            |      |            |                          |              |      |        |                 |             |
| GTGC                      | BAAAAAC<br>             <br>BAAAAAC | 8=8                       | CITIC                                                       | CAAC                             | 1;                   |                            |               |         | F11      |                                                    |       |        |          |                |            |        |            |       |                |            |      |            |                          |              |      |        |                 |             |
| TITE<br>TITE              | GIO                                 | AAACT                     | ince                                                        | )<br>                            | Len                  |                            |               |         | Ø        |                                                    |       |        |          |                |            |        |            |       |                |            |      |            |                          |              |      |        |                 |             |
| 661<br>101                | 300<br>100<br>100                   | HIGG<br>HIGG              | 160                                                         | 8=8                              | lgth<br>Inde         |                            |               |         | Wrap     |                                                    |       |        |          |                |            |        |            |       |                |            |      |            |                          |              |      |        |                 |             |
| TIGG                      | GAG                                 | 160                       | TIG                                                         | 160                              | 19<br>1s             |                            |               |         | per      |                                                    |       |        |          |                |            |        |            |       |                |            |      |            |                          |              |      |        |                 |             |
| 8=8                       | AAG                                 | GAG                       | 6<br>6<br>6<br>6<br>6<br>7<br>6<br>7                        | ACA.                             | 67;                  |                            |               |         | or       |                                                    |       |        |          |                |            |        |            |       |                |            |      |            |                          |              |      |        |                 |             |
| GCAG                      | 6667<br>                            | GAC                       | FIGCT                                                       | AAAGC<br>     <br>!AAAGC         | 0;                   |                            |               |         | PALM     |                                                    |       |        |          |                |            |        |            |       |                |            |      |            |                          |              |      |        |                 |             |
| 3 – 3<br>1                | AG 2<br>AG 1                        | AC 1                      | ล_ล<br>า                                                    | 8=8                              | Gap                  |                            |               |         | ¥        |                                                    |       |        |          |                |            |        |            |       |                |            |      |            |                          |              |      |        |                 |             |
| .648                      | 240<br>588                          | .80<br>.528               | 20<br>468                                                   | 50<br>L408                       | sq                   |                            |               |         |          |                                                    |       |        |          |                |            |        |            |       |                |            |      |            |                          |              |      |        |                 |             |
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                                                                                                                                                                                                                                                                                                                                                                   Sequence 15734, Application US/60172373

GENERAL INFORMATION:
APPLICANT: Morris, MacDonald
APPLICANT: Lal, preet1
APPLICANT: Lolep, Dinh
TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
TITLE OF INVENTION: Polynucleotide Sequence Databases, and Single Nucleotide Polymorp
FILE REFERENCE: GX-0006 p
CURRENT APPLICATION NUMBER: US/60/172,373
CURRENT APPLICATION NUMBER: US/60/172,373
CURRENT FILING DATE: 1999-12-16
NUMBER OF SEQ ID NOS: 25,772
SOFTWARE: PERL PROGram
SEQ ID NO 15734
LENGTH: 716
                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -60-172-373-15734
                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No: 227484.3c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1410
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                                                                                    GACACGAGCAGGAGAGAGAGAGAGAAAAGAAAAACACAGCATGAGAACACAGTAAATG
AGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGATCATG
                                                         CGGCAGCACAGCACCTGCACGAACACCCCGCCGAAACTGCTGCGAGGACACCGTGTACAGG 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCA 366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CGGCAGCACAGCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACACCGTGTACAGG
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                                                                                                                                                                                                                                h 87.2%;
Similarity 99.8%;
                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                             Score 545; DB o.,
Pred. No. 8e-108;
                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                          DB 61; Length 716;
                                                                                                                                                                                                                 Indels
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APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
APPLICANT: Molesh, David Alan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.48401PC
CURRENT APPLICATION UNMBER, PCT/US01/45395
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 215
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LENGTH: 1619;
TYPE: DNA;
ORGANISM: Homo sapiens
PCT-US01-45395-205
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PCT-US01-45395-205
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                                                                                                                                                                                                                                                                                        Matches
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                    1112
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         240 GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTTGGCCGCAG
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                                                                                                                                                                                                                                                                                                  Match 77.8%;
Local Similarity 99.8%;
                                                                                                                                                                                                                                          1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60
                                                                                                                             CAGCGACAGGCGGCAGCACAGCACC-TGCACGAACACCCGGCGAAACTGCTGCGAGGACA 179
                                                                                                GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 1111
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                                                                                                                                                                                                                                                                                      606;
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                                                                                                                                                                                                                                                                                                Score 486; DB 1;
Pred. No. 2.7e-95;
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1232 GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTTGGCCGCAG 1291
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(Sequence 205, Application US/09713550

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(SENERAL INFORMATION:

(APPLICANT: XL, Jiangchun

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(APPLICANT: XL, Jiangchun

(APPLICANT: XL, Jiangchun

(APPLICANTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER

(TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER

(TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CANCER

(TITLE OF INVENTION: 2010-11-14

(CURRENT APPLICATION NUMBER: 2000-11-14

(NUMBER OF SEQ ID NOS: 205

(CURRENT FILING DATE: 2000-11-14
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SEQ ID NO 205
LENGTH: 1619
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ORGANISM: Homo sapiens
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                     1292
                                                                                                                                                           1172 CCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAAACGTCTCCGAGAAGGGGA 1231
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Local Similarity 99.8%;
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                  CCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC 359
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                                                                                                                                                                           CCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGA 239
                                                                                        GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAG
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| 486; | 1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCG | 1532  1592  1592  1592  1592  APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA APPLICA | 540 | Qy 480 TGGTACCAATTTTTCAGTGTTGACAGCTT               | OY 420 GTAAATGAATAAAACCATAAAATATTTAGCCCCTC                           | Qy 360 CTGGGCAGACACGAGCAGGAGGAGAGAGAGACAGAGAA<br> | Qy 300 CCATGATCCTCCGAATCTGGTTGGGCATCCAGCAT                          | Db 1232 GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTC                      |
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|      | 48<br>No<br>sma                      | 95<br>95<br>MET<br>F OV                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | 5 9 | AGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAGAGAGAAATT 539 | GTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA 479<br> | 8 – 8                                             | CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC 359<br> | GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAG 129 |

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Copyright (c) 1993 - 2002
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/cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
/cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*
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US-10-264-237-1044
US-10-264-237-464
US-10-264-237-466
US-09-721-456-3
US-09-513-999C-21763
US-09-513-999C-21763
US-10-131-813A-533
US-10-131-812A-533
US-10-131-826A-533
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           US-10-127-837A-533

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US-10-127-850A-533

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464,
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| C 45              | 44                 | 43                | 42                 | 41                | 40                | . (L)              | . u                | 37                 | 36                 | . u                | 34                 | i Lu               | 3 2                | 3.1                | 30                 | 20                 | 28                 | 27                 |
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| 19                | 19                 | 19                | 19                 | 19                | 19                | 19                 | 19                 | 19                 | 19                 | 19                 | 19                 | 19                 | 19                 | 19                 | 19                 | 19                 | , L                | 19                 |
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| 15832             | 3196               | 2069              | 1330               | 692               | 531               | 496                | 496                | 496                | 496                | 496                | 496                | 496                | 496                | 496                | 496                | 496                | 496                | 496                |
| 6                 | 6                  | σ                 | σ                  | 6                 | 4                 | σ                  | σ                  | σ                  | 6                  | Q                  | 0                  | σ                  | σ                  | σ                  | თ                  | σ                  | 0                  | თ                  |
| US-10-240-453-117 | US-10-264-237-1221 | US-10-264-237-447 | US-10-264-237-1069 | US-10-264-237-662 | US-08-250-795A-19 | US-10-137-873A-533 | US-10-137-872A-533 | US-10-131-836A-533 | US-10-131-821A-533 | US-10-128-693A-533 | US-10-128-690A-533 | US-10-128-686A-533 | US-10-128-685A-533 | US-10-128-684A-533 | US-10-127-849A-533 | US-10-127-843A-533 | US-10-127-838B-533 | US-10-127-825A-533 |
|                   | 1221               | 447               | 1069               | 662               | Sequence 19, Appl | Sequence 533, App  | Sequence 533, App  | 533                | Sequence 533, App  | 533                | Sequence 533, App  | 533                | Sequence 533, App  | Sequence 533, App  | Sequence 533, App  | Sequence 533, App  | Sequence 533, App  | Sequence 533, App  |

ALIGNMENTS

## APPLICANT: Beker, Kevin P. APPLICANT: Desnoyers, Luc APPLICANT: Gedritsen, Mary APPLICANT: Gedritsen, Mary APPLICANT: Goddard, Audrey APPLICANT: Goddard, Audrey APPLICANT: Grimaldi, J. Christopher APPLICANT: Grimaldi, J. Christopher APPLICANT: Godowski, Paul J. APPLICANT: Godowski, Paul J. APPLICANT: Smith, Victoria APPLICANT: Smith, Victoria APPLICANT: Smith, Victoria APPLICANT: Watanabe, Colin L. APPLICANT: Watanabe, Colin L. APPLICANT: Watanabe, Colin L. APPLICANT: Watanabe, Colin L. APPLICANT: Watanabe, Colin L. APPLICANT: Watanabe, Colin L. APPLICANT: Watanabe, SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3530P1C94 CURRENT FILING DATE: 19302-08-28 PRIOR APPLICATION NUMBER: US/10/230,437 CURRENT FILING DATE: 1937-09-17 PRIOR APPLICATION NUMBER: 60/06213 PRIOR APPLICATION NUMBER: 60/06287 PRIOR APPLICATION NUMBER: 60/06287 PRIOR APPLICATION NUMBER: 60/064103 PRIOR FILING DATE: 1937-10-31 PRIOR APPLICATION NUMBER: 60/064103 PRIOR APPLICATION NUMBER: 60/069873 PRIOR APPLICATION NUMBER: 60/078910 PRIOR APPLICATION NUMBER: 60/079910 PRIOR APPLICATION NUMBER: 60/079294 PRIOR APPLICATION NUMBER: 60/079294 PRIOR APPLICATION NUMBER: 60/079294 PRIOR APPLICATION NUMBER: 60/079294 PRIOR APPLICATION NUMBER: 60/079294 PRIOR APPLICATION NUMBER: 60/079294 ; TYPE: DNA ; ORGANISM: Homo US-10-230-437-15 RESULT 1 US-10-230-437-15 Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 246 SEQ ID NO 15 LENGTH: 1524 Sequence 15, Application US/10230437 GENERAL INFORMATION: PRIOR APPLICATION NUMBER: 60/079728 PRIOR FILING DATE: 1998-03-27 PRIOR APPLICATION NUMBER: 60/ PRIOR FILING DATE: 1998-03-26 60/079656

Sapien

62.9%;

Score 393;

В 6,

Length 1524;

Matches

1132

1192

61

1252

181

121

Best Local Similarity

Conservative

100.0%;

Pred. No. 9.3e-177;

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; NAME/KEY: misc_feature; LOCATION: (1297)..(1297)
; OTHER INFORMATION: n equals a,t,g, or US-10-264-237-1044
Sequence 1380, Application US/10264237
GENERAL INFORMATION:
APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PA131P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1044, Application US/10264237 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 2876
SOFTWARE: PATENTIN Ver. 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PA131P1
CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
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                                         ; NAME/KEY: misc_feature
; LOCATION: (1998)..(1998)
; OTHER INFORMATION: n equals
US-10-264-237-464
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US-10-264-237-464
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SEQ ID NO 1380
LENGTH: 2049
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 464, Application US/10264237 GENERAL INFORMATION:
                                                                                                                                                                                                   SOFTWARE: Pa
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CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILLING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 2876
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SOFTWARF: Details
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA131P1
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OTHER INFORMATION: n equals a,t,g,
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                                                                                                                                       ORGANISM: Homo sapiens
                                                                                                                                                                TYPE: DNA
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LOCATION: (1987)..(198
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             325 GTGNAAAAAAAAAAAAAAAA 304
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                                                                                                                                                                                   2062
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Pred. No.
        Score 21;
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          Length 2062;
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US-10-264-237-1044

SEQ ID NO 1044

LENGTH: 1329

FEATURE:

ORGANISM: Homo sapiens

RESULT 3 US-10-264-237-1380/c

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Matches Best Query Match

Local

Similarity

Conservative

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DB 5; 0.29;

Length 2176;

Indels

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US-09-721-456-3
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                                                                                  TELEFAX: (415) 576-0 INFORMATION FOR SEQ ID NO: /no SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1996 TGNAAAAAAAAAAAAAAAA 2016
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                                                                                                                                          TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/721,456
FILING DATE: 22-Nov-2000
                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Human Telomerase Catalytic Subunit NUMBER OF SEQUENCES: 727
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                    LENGTH: 2176 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                               NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
                                                                                                                                                                                                                                                              APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/974,549A
FILING DATE: 19 NOV-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 08/851,843 FILING DATE: 06-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/846,017 FILING DATE: 25-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US.08/844,419 FILING DATE: 18-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: California
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                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/915,503 FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/912,951 FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/911,312 FILING DATE: 14-AUG-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/854,050 FILING DATE: 09-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Townsend and
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Harley, Calvin B.
Andrews, William H.
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Chapman, Karen B.
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                                                                                                                         576-0300
                                                                                                     /note= "clone 712562"
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; OTHER INFORMATION: Genbank Accession No. AA631399
US-10-240-425-208
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SOFTWARE: PatentIn Ver. 2.
SEQ ID NO 208
                                   SOFTWARE: Patent.pm
SEQ ID NO 21763
LENGTH: 163
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Best Local Similarity
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Best Local
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TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.

FILE REFERENCE: 59.US2.REG
                                                                                          PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
                                                                                                                                                                                                                                   APPLICANT: APPLICANT:
                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/513,999C CURRENT FILING DATE: 2000-02-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/240,425
CURRENT FILING DATE: 2002-09-30
PRIOR APPLICATION NUMBER: PCT/US01/09847
PRIOR FILING DATE: 2001-03-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Wetzel, Jon C.
APPLICANT: Scherf, Uwe
APPLICANT: Vockley, Joseph G.
TITLE OF INVENTION: Gene Expression Profiles in Esophageal Tissue
FILE REFERENCE: 44921-5026
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APPLICANT:
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PRIOR FILING DATE: 2000-03-31
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ORGANISM: Homo sapiens
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LOCATION: 1..2176
SEQUENCE DESCRIPTION: SEQ ID NO:
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Lord, Reginald V.
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Pred. No.
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Pred. No.
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DB 6; 0.91;

Length 457;

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Indels

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; FEATURE:
, NAME/KEY: misc_feature
, LOCATION: 135
, OTHER INFORMATION: n-a,
US-09-513-999C-21763
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                                                                    Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENGTH: 496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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APPLICANT:
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CURRENT APPLICATION NUMBER: US/10/131,813A
CURRENT FILING DATE: 2002-04-24
PRIOR APPLICATION NUMBER: 60/049911
PRIOR PELICATION NUMBER: 60/049911
PRIOR PELICATION NUMBER: 60/049911
PRIOR PELICATION NUMBER: 60/056974
PRIOR PELICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR FILING DATE: 1997-08-26
                                                                                                                                             PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/059113
PRIOR TILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
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                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/059117 PRIOR FILING DATE: 1997-09-17
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                                TYPE: DNA ORGANISM: Homo Sapien
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LOCATION: 134
OTHER_INFORMATION: w=a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
 NAME/KEY: unsure
                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      135 NAAAAAAAAAAAAAAAA 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         607 NAAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                                                         FILING DATE: 1997-09-18
                                                                                                                                                                                                                                          APPLICATION NUMBER:
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Watanabe, Colin K
Wood, William
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Godowski, Paul J.
Gurney, Austin L.
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Stewart, Timothy A
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v 100.0%; Pro
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                                                                                                                                                                                                                                          60/059263
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Pred. No. 2.8;
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                                                                                                                              See File Wrapper or PALM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 163
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                                                     ; FEATURE:
; NAME/KEY: unsure
; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-131-819A-533
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US-10-131-819A-533
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                                                                                                                                                                                   SEQ ID NO 533
LENGTH: 496
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-08-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/131,819A CURRENT FILING DATE: 2007-04-24 PRIOR APPLICATION NUMBER: 60/049911 PRIOR FILING DATE: 1997-06-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C134
                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR
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PRIOR APPLICATION NUMBER: 60/059115
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                                                                                                                                               TYPE: DNA ORGANISM: Homo Sapien
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                                                                                                                                                                                                                                                                                                                 FILING DATE: 1997-09-18 APPLICATION NUMBER: 60/059352
                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/059117 FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Desnoyers, Luc
Filvaroff, Ellen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Godowski, Paul J
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 3.0%;
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L00.0%; Pred. No.
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 Score 19;
Pred. No.
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. 2.7;
 DB 6;
2.7;
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                   Length 496;
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                                                                                                                                                                                                                                         SEQ ID NO 533
LENGTH: 496
TYPE: DNA
                                                                    Matches
                                                                               Query Match
Best Local Similarity
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APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Maur
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PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-19
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
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                                                                                                                                                                                                                                                                                            Remaining Prior Application data removed - See File Wrapper or NUMBER OF SEQ ID NOS: 550
                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1997-09-19
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                                                                                                                                                                     NAME/KEY: unsure
LOCATION: 396
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                   607 NAAAAAAAAAAAAAAAAA 625
396 NAAAAAAAAAAAAAAAA 414
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APPLICATION NUMBER: 60/059184
FILING DATE: 1997-09-17
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Watanabe, Colin K
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                                                                  Conservative
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                                                                                3.0%; Score 19;
100.0%; Pred. No.
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                                                                Mismatches
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RESULT 12
US-10-131-826A-533
; Sequence 533, Application US/10131826A
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
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                                                                                                                                                                                                                                                              ; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-131-824A-533
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US-10-131-824A-533
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/05
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/05
PRIOR FILING DATE: 1997-09-17
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Best Local S
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LENGTH: 496
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PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C126
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PRIOR APPLICATION NUMBER: 60/
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PRIOR APPLICATION WHITE
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ORGANISM: Homo Sapien
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19; Conserv
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: 1997-09-17
NUMBER:
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Pred. No.
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Beresini, Maureen

Desnoyers,Luc DeForge,Laura

Goddard, Audrey

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CURRENT APPLICATION NUMBER: US/10/131,826A
CURRENT FILING DATE: 2002-04-24
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR FILING DATE: 1997-06-674
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/05911
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
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US-10-125-926A-533
: Sequence 533, Application US/10125926A
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C138
CURRENT APPLICATION NUMBER: US/10/131,829A
CURRENT FILING DATE: 2002-04-27
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PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
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APPLICANT:
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PRIOR FILING DATE: 1997-09-19
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ORGANISM: Homo Sapien
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NAME/KEY: unsure
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                                    Sherwood, Steven
                                                        Gurney, Austin L.
                                                                        Goddard, Audrey
Godowski, Paul J.
                                                                                                               Gerritsen, Mary E.
                                                                                                                                   Gao, Wei-Qiang
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Filvaroff, Ellen
                                                                                                                                                                                              DeForge, Laura
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: 1997-09-17
NUMBER:
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100.0%; Pred. No.
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; OTHER INFORMATION: unknown base US-10-131-826A-533

FEATURE: NAME/KEY: unsure LOCATION: 396

SEQ ID NO 533 LENGTH: 496

TYPE: DNA

ORGANISM: Homo Sapien

NUMBER OF SEQ ID NOS: 550

PRIOR APPLICATION NUMBER: 60/059588 PRIOR FILING DATE: 1997-09-19

PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1997-0

1997-09-19

60/059352

PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-18

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APPLICATION NUMBER:

APPLICATION NUMBER: 60/059117 FILING DATE: 1997-09-17

APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT:

Tumas, Daniel Watanabe, Colin K Wood, William Zhang, Zemin

Stewart, Timothy A. Smith, Victoria

Godowski, Paul J. Gurney, Austin L. Sherwood, Steven

Goddard, Audrey Gerritsen, Mary E. Gao, Wei-Qiang Filvaroff, Ellen

Query Match Best Local

Local

h 3.0%; Similarity 100.0%; 19; Conservative

0;

Score 19; pred. No. Mismatches

Matches

19;

RESULT 13 US-10-131-829A-533

Sequence 533, Application US/10131829A GENERAL INFORMATION:

APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Mauree
APPLICANT: DeForge, Laura
APPLICANT: Desnoyers, Luc
APPLICANT: Filvaroff, Eller
APPLICANT: Gao, Wei-Glang
APPLICANT: Gerritsen, Mary

Desnoyers, Luc Filvaroff, Ellen Beresini, Maureen

Tumas, Daniel

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RESULT 15
US-10-127-829A-533
Sequence 533, Application US/10127829A
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Best Local Similarity
Matches 19; Conserv
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Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
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                                      APPLICANT:
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ORGANISM: Homo Sapien
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FILING DATE: 1997-09-18
APPLICATION NUMBER: 60/059352
FILING DATE: 1997-09-19
   CANT: Zhang, Zemin OF INVENTION: SECRI OF INVENTION: ACII
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FILING DATE: 1997-09-17
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FILING DATE: 1997-09-17
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FILING DATE: 1997-09-17
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                                                    Watanabe,Colin K
Wood,William
                                                                                    Stewart, Timothy A. Tumas, Daniel
                                                                                                                                                  Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
                                                                                                                                                                                                     Gerritsen, Mary E
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                                                                                                                        Smith, Victoria
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SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC ACIDS ENCODING THE SAME
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100.0%; Pred. No.
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RESULT 16
US-10-127-831A-533
Carmence 533, Application US/10127831A
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Best Local
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                CURRENT APPLICATION NUMBER: US/10/127,831A CURRENT FILING DATE: 2002-10-15 PRIOR APPLICATION NUMBER: 60/049911 PRIOR FILING DATE: 1997-66-18
                                                                                   TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C107
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
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PRIOR FILING DATE: 1997-06-18
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PRIOR APPLICATION NUMBER: 60/
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PRIOR FILING DATE: 1997-08-26
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RIOR APPLICATION NUMBER: 60/056974
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FILING DATE: 1997-09-17
APPLICATION NUMBER: 60/059122
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Gurney, Austin L.
Sherwood, Steven
                                                                                                                                                    Watanabe,Colin K
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Pred. No.
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DAWN

PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-17

60/059117 60/059115 60/059113

FILING DATE: 1997-0: APPLICATION NUMBER: FILING DATE: 1997-01 APPLICATION NUMBER:

1997-09-17 1997-08-26

PRIOR APPLICATION NUMBER: 60/059122 PRIOR FILING DATE: 1997-09-17

PRIOR FILING DATE: 1997-09-19 PRIOR APPLICATION NUMBER: 60/PRIOR FILING DATE: 1997-09-19

60/059588 60/059352

PRIOR APPLICATION NUMBER:

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Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENGTH: 496
TYPE: DNA
ORGANISM: Homo Sapien
                                                     TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C102
CURRENT APPLICATION NUMBER: US/10/127,835A
CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR APPLICATION NUMBER: 60/056974
PRIOR APPLICATION NUMBER: 60/056974
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100.0%; Pred. No.
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US-10-127-837A-533
; Sequence 533, Application US/10127837A
; GENERAL INFORMATION:
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CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059115
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PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENGTH: 496
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NAME/KEY: unsure
LOCATION: 396
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APPLICATION NUMBER: 60/059122
FILING DATE: 1997-09-17
APPLICATION NUMBER: 60/059184
FILING DATE: 1997-09-17
                                                                                   APPLICATION NUMBER: 60/059117 FILING DATE: 1997-09-17
                                        APPLICATION NUMBER: 60/059122
FILING DATE: 1997-09-17
APPLICATION NUMBER: 60/059184 FILING DATE: 1997-09-17
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Wood, William
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US-10-127-831A-533

NAME/KEY: unsure LOCATION: 396 OTHER INFORMATION: unknown base

FEATURE

Matches Query Match

Local Similarity es 19; Conserva

Conservative

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RESULT 17 US-10-127-835A-533 ; Sequence 533, Application US/10127835A

GENERAL INFORMATION:

Baker, Kevin P

APPLICANT:

APPLICANT:

APPLICANT:

APPLICANT:

Watanabe, Colin K Wood, William 2hang, Zemin

Tumas, Daniel Smith, Victoria

Stewart, Timothy A

APPLICANT:

APPLICANT:

PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17

APPLICATION

NUMBER: 60/059117

APPLICANT:

Gerritsen, Mary E.
Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven

APPLICANT: APPLICANT:

APPLICANT:

Gao, Wei-Qiang Filvaroff, Ellen

Desnoyers, Luc DeForge, Laura Beresini, Maureen

APPLICANT: APPLICANT:

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                                                                                                                                                                                                                                                    CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENGTH: 496
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CURRENT FILING DATE: 2002-10-15
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PRIOR APPLICATION NUMBER: 60//59352
PRIOR FILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
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ORGANISM: Homo Sapien
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LOCATION: 396
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   607 NAAAAAAAAAAAAAAAA 625
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                                                                                                                                                                              APPLICATION NUMBER: 60/
FILING DATE: 1997-09-17
                                                                                                                                                                                                                  APPLICATION NUMBER: 60/
FILING DATE: 1997-09-17
                          APPLICATION NUMBER: 60/059588 FILING DATE: 1997-09-19
                                                                FILING DATE: 1997-09-19
                                                                                  APPLICATION NUMBER: 60/
                                                                                                                      APPLICATION NUMBER: 60/059263
                                                                                                                                        APPLICATION NUMBER: 60/059184 FILING DATE: 1997-09-17
        Prior Application
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Zhang, Zemin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wood, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Watanabe, Colin K
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Stewart, Timothy A.
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100.0%; Pred. No.
tive 0; Mismatc
                                                                                  60/059352
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                                                                                                                                                                                                60/059122
data removed - See File Wrapper or PALM.
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2.7;
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RESULT 20
US-10-127-850A-533
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                                                         NUMBER OF SE
SEQ ID NO 533
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SEQ ID NO 533
LENGTH: 496
TYPE: DNA
ORGANISM: Homo Sapien
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Best Local :
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CURRENT APPLICATION NUMBER: US/10/127,850A

CURRENT FILING DATE: 2002-10-15

PRIOR APPLICATION NUMBER: 60/049911

PRIOR PILING DATE: 1997-06-18

PRIOR PILING DATE: 1997-08-26

PRIOR PILING DATE: 1997-08-26

PRIOR PILING DATE: 1997-08-26
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                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/PRIOR FILING DATE: 1997-09-17
                                                                          NUMBER OF SEQ ID NOS: 550
                                                                                                            PRIOR FILING DATE:
                                                                                                                                               PRIOR APPLICATION NUMBER: 60/059352
                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/059117 PRIOR FILING DATE: 1997-09-17
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TYPE: DNA
ORGANISM: Homo Sapien
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LOCATION: 396
OTHER INFORMATION: unknown base
                                        LENGTH: 496
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                                                                                                                             APPLICATION NUMBER: 60/059588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tumas, Daniel
Watanabe, Colin K
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                                                                                                              1997-09-19
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                                                                                           data removed
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    See File Wrapper or PALM.

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TITLE OF INVENTION: SECRITED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C86
CURRENT APPLICATION NUMBER: US/10/127,901A
CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/05913
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
PRIOR APPLICATION NUMBER: 60/05912
PRIOR APPLICATION NUMBER: 60/05912
PRIOR APPLICATION NUMBER: 60/05912
PRIOR APPLICATION NUMBER: 60/05912
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/05912
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; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-127-850A-533
                                   ; FEATURE:
; NAME/KEY: unsure
; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-127-901A-533
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US-10-127-901A-533
                                                                                                                                                   Remaining Prior Application
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENGTH: 496
TYPE: DNA
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Best Local :
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Query Match
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                                                                                                                               ORGANISM: Homo Sapien
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es 19; Conserv
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Wood, William
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Gurney, Austin L.
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Filvaroff, Ellen
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100.0%; Pr
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   Score 19;
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   DB 6;
                                                                                                                                                                                                                                 See File Wrapper or PALM
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                                         607 NAAAAAAAAAAAAAAAAA 625
     396 NAAAAAAAAAAAAAAAAA 414
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RESULT 22
US-10-128-689A-533
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CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/049911
PRIOR ETLING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059117
PRIOR PRILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
                                                                                    US-10-128-689A-533
                                                                                                                                                                                                                      Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENGTH: 496
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Matches 19; Conserv
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                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/059352 PRIOR FILING DATE: 1997-09-19
                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
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                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 1997-09
PRIOR APPLICATION NUMBER:
                                                                                                   NAME/KEY: unsure
LOCATION: 396
OTHER INFORMATION: unknown base
                                                                                                                                                                TYPE: DNA
ORGANISM: Homo Sapien
FEATURE:
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                         Local Similarity
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Wood,William
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      Conservative
100.08; ++
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                       Score 19;
; Pred. No.
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                                              DB 6;
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                                            Length 496;
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APPLICANT:

Baker, Kevin P.

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US-10-131-833A-533

Sequence 533, Application US/10131833A

GENERAL INFORMATION:
                                                                                                                                                                                                                  : FEATURE:
NAME/KEY: unsure
LOCATION: 396
CHER INFORMATION: unknown base
US-10-131-830A-533
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US-10-131-830A-533
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SEQ ID NO 533
LENGTH: 496
TYPE: DNA
                                                                                                                                                                      Query Match
Best Local s
                                                                                                                                                         Matches
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PRIOR FILING DATE: 1997-09-17
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PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
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                                                                                                         607 NAAAAAAAAAAAAAAAAA 625
                                                                                         396 NAAAAAAAAAAAAAAAA 414
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FILING DATE: 1997-09-19
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FILING DATE: 1997-09-18
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                                                                                                                                                                      Similarity
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Godowski, Paul J.
Gurney, Austin L.
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Wood,William
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1997-09-17
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2.7;
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RESULT 25
US-10-131-837A-533
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CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR PRIOR DATE: 1997-06-26
PRIOR PRIOR DATE: 1997-09-17
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PRIOR PRIOR DATE: 1997-09-17
PRIOR PRIOR DATE: 1997-09-17
                                                                        Sequence 533, Application US/10131837A
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Maureen
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Best Local Similarity
Matches 19; Conserv
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo Sapien
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Gao, Wei-Qiang
                  Desnoyers, Luc
Filvaroff, Ellen
                                                        DeForge, Laura
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Wood, William
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APPLICANT:

Godowski, Paul J. Gurney, Austin L.

Goddard, Audrey Gerritsen, Mary E.

Sherwood, Steven

PPLICANT:

PPLICANT:

APPLICANT: APPLICANT: \PPLICANT: APPLICANT: APPLICANT:

Stewart, Timothy A. Tumas, Daniel

APPLICANT: PPLICANT:

Zhang, Zemin Wood, William Watanabe, Colin K

APPLICANT:

Stewart, Timothy A.

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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REPERENCE: P33301CL31
CURRENT APPLICATION NUMBER: US/10/131,837A
CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/049911
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR APPLICATION NUMBER: 60/056913
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-08-26
PRIOR FILING DATE: 1997-09-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 496;
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US-10-127-825A-533
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LENGTH: 496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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                                                                                                  APPLICANT:
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PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
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  TITLE
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OTHER INFORMATION: unknown base
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APPLICATION NITHERT
OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 1997-09-
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                                    Watanabe, Colin K
Wood, William
                                                                                              Smith, Victoria
Stewart, Timothy A.
                                                                                                                                         Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
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Gao, Wei-Qiang
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                                                                                 Tumas, Daniel
                                                                                                                                                                                                       Goddard, Audrey
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100.0%; Pred. No.
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2.7;
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Best Matches Query Match

Local 19;

Similarity 100.0%;

3.0%; Score 19; 0; Pred.

DB 6;

Conservative

Mismatches No.

US-10-125-930A-533 RESULT 26

Sequence 533, Applic GENERAL INFORMATION

Application US/10125930A

APPLICANT:

Desnoyers, Luc Filvaroff, Ellen

DeForge, Laura Beresini, Maureen

APPLICANT:

APPLICANT: Baker, Kevin P.

APPLICANT APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT:

Goddard, Audrey Godowski, Paul J. Gurney, Austin L. Sherwood, Steven

Gerritsen, Mary E Gao, Wei-Qiang ; NAME/KEY: unsure ; LOCATION: 396 ; OTHER INFORMATION: unknown base US-10-131-837A-533

SEQ ID NO 533 LENGTH: 496

TYPE: DNA

FEATURE:

ORGANISM: Homo Sapien

PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19 PRIOR APPLICATION NUMBER: 60/059588 PRIOR FILING DATE: 1997-09-19

PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059263

PRIOR APPLICATION NUMBER: 60/059184

PRIOR APPLICATION NUMBER: 60/059122 PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/059115 PRIOR FILING DATE: 1997-09-17

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; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-127-825A-533
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
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CURRENT APPLICATION NUMBER: US/10/127,838B
CURRENT FILING DATE: 2002-04-22
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
                                                                    APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C98
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PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
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PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
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CURRENT APPLICATION NUMBER: US/10/127,825A
CURRENT FILING DATE: 2002-04-22
PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
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                                                                                                                                                               Tumas, Daniel
Watanabe, Colin K
Wood, William
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Godowski, Paul J.
Gurney, Austin L.
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CURRENT APPLICATION NUMBER: US/10/127,843A CURRENT FILING DATE: 2002-04-22 PRIOR APPLICATION NUMBER: 60/049911 PRIOR FILING DATE: 1997-06-18 PRIOR PLICATION NUMBER: 60/056974 PRIOR ETLING DATE: 1997-08-26 PRIOR ETLING DATE: 1997-09-17 PRIOR ETLING DATE: 1997-09-17 PRIOR ETLING DATE: 1997-99-17
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PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
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PRIOR FILING DATE: 1997-08-26
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NAME/KEY: unsure
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APPLICATION NUMBER: 60/059115 FILING DATE: 1997-09-17
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APPLICATION NUMBER: 60/059263
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        533, Application US/10127843A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Baker, Kevin P.
                                                                                                                                                                                                                                           Zhang, Zemin
                                                                                                                                                                                                                                                            Watanabe,Colin K
Wood,William
                                                                                                                                                                                                                                                                                                 Tumas,Daniel
                                                                                                                                                                                                                                                                                                                                                                    Godowski, Paul J.
Gurney, Austin L.
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Filvaroff, Ellen
                                                                                                                                                                                                                                                                                                              Stewart, Timothy A.
                                                                                                                                                                                                                                                                                                                                   Smith, Victoria
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18

PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17

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PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
PREMAINING PRIOR APPLICATION data removed - See File Wrapper or PALM.
NUMBER OF SEO ID NOS: 550
SEO ID NO 533
LENGTH: 496
                                                                                                                                        TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R10103

CURRENT APPLICATION NUMBER: US/10/127,849A

CURRENT FILING DATE: 2002-04-23

PRIOR APPLICATION NUMBER: 60/049911

PRIOR APPLICATION NUMBER: 60/049911

PRIOR APPLICATION NUMBER: 60/05974

PRIOR APPLICATION NUMBER: 60/05974

PRIOR APPLICATION NUMBER: 60/059113

PRIOR FILING DATE: 1997-08-26

PRIOR FILING DATE: 1997-09-17

PRIOR FILING DATE: 1997-09-17
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2.7;
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; OTHER INFORMATION: unknown base
US-10-127-849A-533
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US-10-128-684A-533
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         PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
PRIOR FILING DATE: 1997-09-19
                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR PELLING DATE: 1997-08-26
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059117
PRIOR APPLICATION NUMBER: 60/059117
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PRIOR FILING DATE: 1997-09-17
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
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PRIOR EILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
PRIOR EILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
PRIOR FILING DATE: 1997-09-19
                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C118
CURRENT APPLICATION NUMBER: US/10/128,684A
CURRENT FILING DATE: 2002-04-23
CURRENT FILING DATE: 2002-04-23
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APPLICANT:
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NAME/KEY: unsure
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           607 NAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match 3.0%; Score 19; Local Similarity 100.0%; Pred. No.
FILING DATE: 1997-09-19
APPLICATION NUMBER: 60/059588
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RESULT 30 US-10-127-849A-533

Sequence 533, Application US/10127849A GENERAL INFORMATION:

APPLICANT: Baker,Kevin P.

APPLICANT:

APPLICANT:

Deforge, Laura Beresini, Maureen Filvaroff, Ellen Desnoyers,Luc

APPLICANT:

Goddard, Audrey Godowski, Paul J. Gurney, Austin L.

Gerritsen, Mary E. Gao, Wei-Qiang

APPLICANT:

APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT:

APPLICANT:

Tumas, Daniel Stewart, Timothy A. Smith, Victoria Sherwood, Steven

APPLICANT: APPLICANT:

Watanabe, Colin K Wood, William

Zemin

PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/059117

FILING DATE: 1997-09-17 APPLICATION NUMBER: 60/059122 FILING DATE: 1997-09-17

APPLICATION NUMBER: 60/ FILING DATE: 1997-09-17

60/059184

US-10-127-843A-533

NAME/KEY: unsure LOCATION: 396

OTHER INFORMATION: unknown base

FEATURE:

TYPE: DNA ORGANISM: Homo Sapien

Query Match Best Local (

Local Similarity

3.0%; Score 19; 100.0%; Pred. No. tive 0; Mismatc

Conservative

Matches

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US-10-128-684A-533
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US-10-128-685A-533
Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENCTH: 496
TYPE: DNA
ORGANISM: Homo Sapien
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SEQ ID NO 533
LENGTH: 496
TYPE: DNA
ORGANISM: Homo Sapien
FEATURE:
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PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-19
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059184
PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P333071C1116
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                                                                                                            PRIOR APPLICATION NUMBER: 60/059588 PRIOR FILING DATE: 1997-09-19
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PRIOR ETLING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
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LOCATION: 396
OTHER INFORMATION: unknown base
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FILING DATE: 1997-09-17
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Godowski, Paul J.
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100.0%; Pred. No.
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                                                                                  See File Wrapper or PALM
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; NAME/KEY: unsure
; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-128-686A-533
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; NAME/KEY: unsure
; LOCATION: 396
; OTHER INFORMATION: unknown base
US-10-128-685A-533
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                                                                                                                                                            SEQ ID NO 533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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Best Local Similarity 100.0%;
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                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/059263
PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
PRIOR APPLICATION NUMBER: 60/059588
PRIOR FILING DATE: 1997-09-19
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PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C119
CURRENT APPLICATION NUMBER: US/10/128,686A
CURRENT FILING DATE: 2002-04-23
CURRENT FILING DATE: 2002-04-23
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
                                                                                              TYPE: DNA
ORGANISM: Homo Sapien
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                                                                                 FEATURE:
                                                                                                                                           LENGTH: 496
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FILING DATE: 1997-09-17
APPLICATION NUMBER: 60/059184
FILING DATE: 1997-09-17
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Filvaroff, Ellen
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Wood,William
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Godowski, Paul J.
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100.0%; Pred. No.
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2.7;
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RESULT 34
US-10-128-690A-533
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                                                                                     Query Match
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LENGTH: 496
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PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
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PRIOR FILING DATE: 1997-09-17
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                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/059352
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                                                                                                                                          NAME/KEY: unsure
LOCATION: 396
OTHER INFORMATION: unknown base
                                                                                                                                                                                                               ORGANISM: Homo Sapien
                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                FEATURE:
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                  607 NAAAAAAAAAAAAAAAAA 625
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                                                  Local Similarity es 19; Conserv
                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 1997-09-19
Wood, William
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Godowski, Paul J.
Gurney, Austin L.
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RESULT 35
US-10-128-693A-533
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RESULT 36
US-10-131-821A-533
; Sequence 533, Application US/10131821A
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PRIOR FILING DATE: 1997-06-18
PRIOR FILING DATE: 1997-08-26
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
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                                                                                                                                                                                                                                                         LOCATION: 396
OTHER INFORMATION: unknown base
                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                                                                                             NAME/KEY: unsure
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                                                                                                                607 NAAAAAAAAAAAAAAAAAA 625
                                                                                                                                                                  Local Similarity hes 19; Conserv
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FILING DATE: 1997-09-19
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                                                                                           NAAAAAAAAAAAAAAA 414
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                                                                                                                                                                  Conservative
                                                                                                                                                                                      100.0%;
                                                                                                                                                                                     3.0%; Score 19;
L00.0%; Pred. No.
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                                                                                                                                                                     0;
                                                                                                                                                                     Mismatches
                                                                                                                                                                                      DB 6;
2.7;
                                                                                                                                                                     0,
                                                                                                                                                                                                          Length 496;
                                                                                                                                                                       Indels
                                                                                                                                                                       0;
                                                                                                                                                                       Gaps
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INFORMATION:

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RESULT 37
US-10-131-836A-533
; Sequence 533, Application US/10131836A
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin P.
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PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 533
LENGTH: 496
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APPLICANT:
                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 1997-09-19
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                                                                                                                                                                                                                                                                                                                                   NAME/KEY: unsure
LOCATION: 396
OTHER INFORMATION: unknown base
                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo Sapien
                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                       607 NAAAAAAAAAAAAAAAA 625
                                                                                                                                                                                   396 NAAAAAAAAAAAAAAAA 414
                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 60/059122 FILING DATE: 1997-09-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
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                              Beresini, Maureen
DeForge, Laura
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Wood, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tumas,Daniel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Smith, Victoria
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Godowski, Paul J.
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                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application data removed - See File Wrapper or PALM NOS: 550
                                                                                                                                                                                                                                                                      3.0%; Score 19; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                   Length 496;
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                                                                                                                                                                                                                                                     Gaps
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Sequence 533, Applic GENERAL INFORMATION:

Application US/10137872A

APPLICANT: Baker, Kevin P. APPLICANT: Beresini, Maur

APPLICANT: APPLICANT: APPLICANT:

APPLICANT:

APPLICANT:

Goddard, Audrey Godowski, Paul J Gurney, Austin L. Sherwood, Steven APPLICANT:

Desnoyers, Luc Filvaroff, Ellen

Beresini, Maureen DeForge, Laura

Gao, Wei-Qiang Gerritsen, Mary E. APPLICANT:

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RESULT 38
US-10-137-872A-533
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PRIOR APPLICATION NUMBER: 60/049911
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR APPLICATION NUMBER: 60/059115
                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 533
LENGTH: 496
                                                                                                                                         Matches
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PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: 1997-0
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PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1997-09-19
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NUMBER OF SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: P3330R1C135
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                                                                                                                                                                                                                        NAME/KEY: unsure
LOCATION: 396
OTHER INFORMATION: unknown base
                                                                                                                                                                                                                                                                                   ORGANISM: Homo Sapien FEATURE:
                                                                                                                                                                                                                                                                                                                        TYPE: DNA
                                                                                        607 NAAAAAAAAAAAAAAAAA 625
                                                                   396 NAAAAAAAAAAAAAAAAA 414
                                                                                                                                           Local Similarity
nes 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 60/059122 FILING DATE: 1997-09-17
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Gerritsen, Mary E.
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Wood, William
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Godowski, Paul J.
                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                          Application data removed - See File Wrapper or PALM
                                                                                                                                                                                                                                                                                                                                                                           NOS:
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                                                                                                                                                          3.0%; Score 19;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                60/059352
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                                                                                                                                           0;
                                                                                                                                           Mismatches
                                                                                                                                                          DB 6;
2.7;
                                                                                                                                                                        Length 496;
                                                                                                                                         Indels
                                                                                                                                         0;
                                                                                                                                         Gaps
                                                                                                                                       0
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Tumas, Daniel

Wood, William Watanabe, Colin K

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; OTHER INFORMATION: unknown base US-10-137-872A-533
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PRIOR FILING DATE: 1997-06-18
PRIOR PELICATION NUMBER: 60/056974
PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 533
LENGTH: 496
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 533, Application US/10137873A
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PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/059352 FILING DATE: 1997-09-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 60/059184 FILING DATE: 1997-09-17
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100.0%; Pred. No.
tive 0; Mismatc
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PRIOR FILING DATE: 1997-08-26
PRIOR APPLICATION NUMBER: 60/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059115
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/059117
PRIOR FILING DATE: 1997-09-17
PRIOR FILING DATE: 1997-09-17
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Best Local
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LENGTH: 496
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PRIOR APPLICATION NUMBER: 60/059352
PRIOR FILING DATE: 1997-09-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 1997-09-19 Remaining Prior Application d NUMBER OF SEQ ID NOS: 550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/059122
PRIOR FILING DATE: 1997-09-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Homo Sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: unsure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: 396
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       396 NAAAAAAAAAAAAAAA 414
                                                               COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII (text)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/250,795A
FILING DATE: 27-May-1994
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Vincent, Matthew
REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                 ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: ROPES & GRAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Immunosuppressant Target Proteins NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Berlin, Vivian
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19; Conserv
                                                                                                                                                                                                                                                                                                                                STATE: MA
                                                                                                                                                                                                                                                                                                                                                    STREET: 1 International Place CITY: Boston
                                                                                                                                                                                                                                                                                                     COUNTRY: USA
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100.0%; Pred. No.
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607

NAAAAAAAAAAAAAAAA 625

Query Match Best Local S Matches 19

Local Similarity

Conservative

NAME/KEY: unsure LOCATION: 396

FEATURE:

ORGANISM: Homo Sapien

PRIOR

PRIOR FILING DATE: 1997-09-19

PRIOR FILING DATE: 1997-09-18 PRIOR APPLICATION NUMBER: 60/

В

396 NAAAAAAAAAAAAAAAAAA 414

RESULT 39 US-10-137-873A-533

GENERAL INFORMATION:

Baker, Kevin P.

Beresini,Maureen

APPLICANT:

Stewart, Timothy A. Tumas, Daniel Watanabe, Colin K Wood, William

APPLICANT APPLICANT APPLICANT: APPLICANT: APPLICANT:

APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT:

> Gerritsen, Mary E. Gao, Wei-Qiang Desnoyers, Luc Filvaroff, Ellen DeForge, Laura

Goddard, Audrey Godowski, Paul J. Gurney, Austin L. Sherwood, Steven

Smith, Victoria

APPLICANT:

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GENERAL INFORMATION:
APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA131P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 2876
SOFTWARE: PATENTIN VET. 3.1
SEQ ID NO 662
ELENCTH: 692
ELENCTH: 692
                                                                 Matches
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Best Local Similarity
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                                                                                                                                       NAME/KEY: misc_feature
LOCATION: (692)..(692)
OTHER INFORMATION: n equals a,t,g, or
                                                                                                                                                                                                       NAME/KEY: misc_feature
LOCATION: (684)..(684)
OTHER INFORMATION: n equals a,t,g, or
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LOCATION: (23)..(23)
OTHER INFORMATION: n equals a,t,g, or c
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OTHER INFORMATION: n equals
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651 NAAAAAAAAAAAAAAAA 669
                             607 NAAAAAAAAAAAAAAAA 625
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LENGTH: 531 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: APBI-P01-036
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 951-7000
TELEPAX: (617) 951-7050
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19; Conserv
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ilarity 100.0%;
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Pred. No.
                                                             Mismatches
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RESULT 44
US-10-264-237-1221
; Sequence 1221, Appli
; GENERAL INFORMATION:
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NAME/KEY: misc_feature

; LOCATION: (2046)..(2046)

; OTHER INFORMATION: n equals a,t,g, or

US-10-264-237-447
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US-10-264-237-447
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NUMBER OF SEQ ID NOS: 2876
SOFTWARE: PATENTIN Ver. 3.1
SEQ ID NO 1069
LENGTH: 1330
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GENERAL INFORMATION:
APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA131P1
                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 3.1
SEQ ID NO 447
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Best Local (
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CURRENT FILLING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: PCT/US01/16450 PRIOR FILING DATE: 2001-05-18
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TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PA131P1
                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR EILING DATE: 2000-05-19
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LOCATION: (1265)...(1265)
OTHER INFORMATION: n equals a,t,g, or
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ORGANISM: Homo sapiens
FEATURE:
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19; Conserv
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Pred. No.
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APPLICANT: Birse et al.

Application US/10264237

and Antibodies

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PRIOR APPLICATION NUMBER: DE 10019058.8
PRIOR FILING DATE: 2000-04-06
PRIOR APPLICATION NUMBER: DE 10019173.8
PRIOR FILING DATE: 2000-04-07
PRIOR ETLING DATE: 2000-04-07
PRIOR ETLING DATE: 2000-06-30
PRIOR ETLING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR APPLICATION NUMBER: DE 10043826.1
PRIOR FILING DATE: 2000-09-01
NUMBER OF SEQ ID NOS: 350
SEQ ID NO 117
LENGTH: 15832
                                , LUCATION: (5221, 5244, US-10-240-453-117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-240-453-117/c
US-10-240-453-117/c
; Sequence 117, Application US/10240453
; GENERAL INFORMATION:
; APPLICANT: OLEK, Alexander
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  Query Match
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SEQ ID NO 1221
LENGTH: 3196
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Diagnosis of Diseases Associated with DNA
TITLE OF INVENTION: Transcription
TITLE OF INVENTION: by Means of Assessing the Methylation Status of Genes Associated
TITLE OF INVENTION: With DNA Transcription
FILE REFERENCE: 5013.1009
CURRENT APPLICATION NUMBER: US/10/240,453
CURRENT FILING DATE: 2002-10-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/264,237
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/16450
PRIOR FILING DATE: 2001-05-18
PRIOR APPLICATION NUMBER: US 60/205,515
PRIOR FILING DATE: 2000-05-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: PCT/EP01/03973 PRIOR FILING DATE: 2001-04-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 2876
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LOCATION: (3170)..(3170)
OTHER INFORMATION: n equals a,t,g,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (668)..(668)
OTHER INFORMATION: n equals a,t,g,
                                                                NAME/KEY: unsure LOCATION: (5221,
                                                                                                                                                                                                OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
                                                                                                                                  NAME/KEY: unsure
LOCATION: (791, 2400,
                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                    TYPE: DNA
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                                                                6508, 12917, 12968, 12987..12988, 13055)
  3.0%;
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Pred. No.
Score 19;
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DB 6;
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Length 15832;
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                        Best Local Similarity 100 Matches 19; Conservative
100.0%; ٢.
                               Pred. No.
                        Mismatches
                              2.3;
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Search completed: November Job time: 48.4612 secs

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Result
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AI936826 wp69h10.x
BE439382 nab63906.
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| 45       | 44                  | 43       | 42       | 41       | 40       | 39       | 38       | 37       | 36                 | 35       | 34       | S<br>S   | 32                 | 31                 | 30                 | 29                 | 28                 | 27                 | 26       | 25                 | 24       | 23       | 22       | 21       | 20                 | 19                 | 18                 | 17       | 16          | 15                 | 14       | 13        | 12       | 11       | 10                                      | 9        | ∞         | 7                  |
| 272      | 282                 | 294      | 298      | 307      | 307      | 312      | 314      | 314      | 314                | 314      | 315      | 332      | 338                | 339                | 339                | 347                | 365                | 368                | 369      | 380                | 390      | 390      | 393      | 396      | 415                | 421                | 423                | 425      | 425         | 437                | 442      | 459       | 462      | 471      | 489                                     | 499      | 516       | 525                |
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| AW128849 | BQ689771            | AI363261 | AI566797 | BG057775 | AA173739 | BF726644 | AA084249 | AA075710 | AA970361           | AA665640 | AI423162 | AW191974 | AI272281           | BF125134           | BF126050           | AW087372           | AI742092           | BF054680           | BF054837 | AI765236           | BE350014 | AI499630 | AI391683 | AA613995 | AI018769           | вм667957           | AI150931           | BE385990 | AI884686    | AW075598           | AW149665 | AI336858  | AW338938 | BF726459 | BF939693                                | BE858216 | AW590950  | BM547680           |
| 28849    | BO689771 AGENCOTIRT | 2        | 70       | 577      | 373      | 266      | 424      | 7571     | AA970361 op52q06.s | 6564     | 2316     | AW1919   | AI272281 ap23c03.x | BF125134 601762356 | BF126050 601762888 | AW087372 xh19g02 x | AI742092 wg38h03 x | BF054680 7169f08 v | BF0548   | AI765236 W172h08 x | BE3500   | 9963     | 9168     | 1399     | AI018769 ov32e04.x | BM667957 UI-E-DW0- | AI150931 ab53c04 x | 3859     | . 705£404 . | AW075598 xh24e07 x | xf40h05  | ax86a11.x | ha68h0   | bv06h03  | 0.0000000000000000000000000000000000000 | 7019007  | ) ha51e13 | BM547680 AGENCOTOT |

RESULT 1
BQ006545/c
LOCUS VERSION KEYWORDS ACCESSION DEFINITION BQ006545 702 bp mRNA linear EST 26-MAR-UI-H-EI1-aza-n-21-0-UI.sl NCI\_CGAP\_EI1 Homo sapiens cDNA clone IMAGE:5846228 3', mRNA sequence. BQ006545.1 GI:19731445 EST 26-MAR-2002

ALIGNMENTS

REFERENCE AUTHORS ORGANISM Homo sapiens numan.

EST

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (Dases 1 to 702)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.

JOURNAL

TITLE

COMMENT

Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tlssue Procurement: Dr. Jose Mercuende
CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Clone distribution information can be found
through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
POLYA-Yes.

Location/Qualifiers
1. .702

FEATURES

source

/clone="IMAGE:5846228" /clone\_lib="NCI\_CGAP\_EI1" /tissue\_type="Chondrosarcoma"

/organism="Homo sapiens" /db\_xref="taxon:9606"

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445 CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60
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                                                                                                 TGGGCAGACACGAGCAGGAGGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note-"Organ: Left Pelvis: Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site_1: EcoR I; Site_2: Not NCI_CGAP_EII is a normalized cDMA library containing the following tissue(s): Chondrosarcoma. The library was
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/lab_host="DH10B (Life Technologies)"
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                                                                                                                                                                                                                                  GGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAG 129
                                                                     GCGGCAGCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAG 418
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1 (bases 1 to 678)

NCI/NIMOS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project
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Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EST
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Seq primer: -40UP from Gibco
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Mammalia; Eutheria; Primates;
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               /note=Porgan: pooled; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from five normalized libraries were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified CDNAs from pools of 5,000 clones made from the same 5 libraries. The pools consisted of the following libraries and clones Soares NbHSF pool 1: 309384-310919, 323208-325895 Soares NbHSF pool 1: 145032-147353, 147730-148103, 148872-149255, 15002 - 150407, 151176-152327 Soares NbZHP-9W pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 Soares NbZHPA pool 1: 758280-760583, 772104-774407 So
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/clone_1ib="Soares_NSF_F8_9W_OT_PA_P_S1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1998)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 537)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index
                                                                                                                                                                                          BF594242
                                                                                                                                                                                                                   BF594242 537 bp mRNA 7n10d03.x1 NCI_CGAP_Brn23 Homo sapiens cisimilar to SW:GP39_HUMAN 043194 PUTATIVE
Tissue Procurement: David
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Soares and M. Fatima Bonaldo."
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Pred. No. 0;
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FEATURES

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CDNA Library Arrayed by: Greg Lennon, Ph.D.

CDNA Sequencing by: Washington University Genome Sequencing Center

DNA Sequencing NCI-CGAP clone distribution information can be
 AI990500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    found through the I.M.A.G.E. Consortium/LLNL, send email to: info@image.llnl.gov
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/clone="IMAGE:3564316"
/clone_lib="NCI_CGAP_Brn23"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /tissue_type="glioblastoma (pooled)"
/lab_host="DH10B"
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                                                                                      AGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGG
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BASE CO

COUNT

Query Match Best Local

70

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AI990500
                                                                     ws40d07.x1 NCI_CGAP_GC6 Homo sapiens cDNA clone IMAGE:2499661 3' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
AI990500.1 GI:5837381
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Homo sapiens

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. (bases 1 to 616)

National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Tumor Gene Index

Unpublished (1997) Contact: Robert Strausberg, Ph.D.

R. Emmert-Buck, M.D., Ph.D. cDNA Library Preparation: M. Bento Soares, Ph.D., Email: cgapbs-r@mail.nih.gov Tissue Procurement: Christopher A. Moskaluk, M.D., M. Fatima Ph.D., Michael

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Cente
Clone distribution: NCI-CGAP clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at: Bonaldo, Ph.D. Genome Sequencing Center əq

High quality sequence stop: Location/Qualifiers 459.

/note="Vector: p77T3D-Pac (Pharmacia) with a modified polylinker; Site\_1: Not I; Site\_2: Eco RI; plasmid DNA from the normalized ilbrary NCI\_CGAP\_GC4 was prepared, and specification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified CDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1257096-1258631, 1469064-1470983, and 147592-1476743). Subtraction by Bento Soares and M. Fatima Bonaldo. "

subtraction by Bento Soares and M. Fatima Bonaldo." /clone="IMAGE:2499661" /clone\_11b="NCI\_CGAP\_GC6" /tissue\_type="pooled germ cell tumors" /lab\_host="DH10B" /organism="Homo sapiens"
/db\_xref="taxon:9606"

Similarity Conservative 84.6%; 0 Score 529; Pred. No. 0; Mismatches DB 9; Length 616; 0, 0;

27 GGGACGCGAAGAGCAACGGGCGCTGCACAAAGCGGGCGCTGTCGGTGGTGGAGTGCGCAT 86 

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GCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCG GTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGGCAGCACCACCCT 146 206

AGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGG GCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCG 266 402

ACCTCGTCCAGTCGTGCTTGGGTTTGGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCA

342

ACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCA

þ mRNA linear EST 08-MAR-2000

RESULT 5 AI990500/c

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FEATURES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    contact: Soares, MB
Program for Rat Gene Discovery and Mapping
University of Iowa
451 Eckstein Moddan
                                                                                                                                                                                                                                                                                                                                                            Email: msoares@blue.weeg.uiowa.edu
Tissue Procurement: Dr. Gregg Hageman
cDNA Library preparation: Dr. M. Bento Soares, Univeristy of Towe
cDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Towe
DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Towa
Clone Distribution: Researchers may obtain clones from Research
Genetics (www.resgen.com).
Seg primer: M13 Forward
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1 (bases 1 to 627)

Bonaldo,M.F., Lennon,G. and Soares,M.B.

Normalization and subtraction: two approaches to facilitate gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     319 335 8250
319 335 9565
/lab_host="DHIOB (Life Technologies) (T1 phage resistant)"
/lab_host="DHIOB (Life Technologies) (T1 phage resistant)"
/note="Organ: eye; Vector: pT7T3-Pac (Pharmacia) with a
modified polylinker; Site_1: EcoR I; Site_2: Not I;
UI-E-DMI is a normalized cDNA library containing the
following tissue(s): lens. The library was constructed
according to Bonaldo, Lennon and Soares, Genome Research,
6:791-806, 1996. First strand cDNA synthesis was primed
with an oligo-dT primer containing a Not I site. Double
stranded cDNA was ligated to an EcoR I adaptor, digested
with Not I, and cloned directionally into pT7T3-Pac
vector. The oligonucleotide used to prime the synthesis of
                                                                                                                                                                                                   /tissue_type="lens"
/dev_stage="adult"
                                                                                                                                                                                                                                       /clone="UI-E-DW1-ahd-c-11-0-UI"
/clone_lib="UI-E-DW1"
                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                             Univeristy of Iowa
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     Contact: Robert
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570 TGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG 606
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                                                                                                                                                           CCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGTTGGACTTGAC
                                                                                                                                                                                                                                        GAGAAAAGAAAAACACAGCATGAGAACACAGTAAATGAATAAAACCATAAAATATTTAGC
                                                                                                                                                                                                                                                                                                                                                AGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAGGAGGAGGAGAGACA
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TGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG
                                                                             AGCTTCTTTTGCCACAAGCAAGAGAGAATTTAACACTGTTTCAAACCCGGGGGAGTTGGC
                                                                                                                                                                                                      CCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGTTGGACTTGAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     first-strand cDNA contains a library tag sequence that is located between the Not I site and the (dT)18 tail. The sequence tag for this library is CGATTAGCGA. This library was created for the program, Gene Discovery in the Visual System, supported by National Eye Institute (NEI). TAG_TISSUE-human lens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   _SEQ=CGATTAGCGA"
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Pred. No. 0;
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BM547680
AGENCOURT\_6507108 NIH\_MGC\_124 Homo sapiens cDNA clone IMAGE:5727798
SION BM547680
BM547680
BM547680.1 GI:18781656
EST.
ELT.
BLARTYOUR HOMO Sapiens
ELMARYOUR HOMO SAPIENS
ELMARYOUR HOMO SAPIENS
ELMARYOUR HOMO
ELMARYOUR HOMO
MAMMALIA; Eutheria; Primates; Catarrhin1; Hominidae; Homo.
1 (bases 1 to 696)
HORS
NIH-MGC http://mgc.nci.nih.gov/.
NH-MGC http://mgc.nci.nih.gov/.
NH-MGC http://mgc.nci.nih.gov/.
NH-MGC http://mgc.nci.nih.gov/.
NH-MGC http://mgc.nci.nih.gov/.
Contact: Robert Strausberg, Ph.D.

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BASE COUNT
ORIGIN
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                                         541 AACACTGTTTCAAACCCGGGGGAGTTGGCTGTGTTA 576
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                                                                                                                                              TGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAG 420
                    AACACTGTTTCAAACCCGGGGGGGTTGGCTGTTTA 640
                                                                                                         TAAATAAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 544
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Tissue Procurement: Invitrogen
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://lmage.llni.gov
Plate: LLAM12722 row: h column: 07
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5727798"
/clone_lib="NIH_MGC_124"
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/lab_host="DH10B"
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99.8%;
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Pred. No. 0;
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TITLE
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100 CTTCTCGTGGTTGGCGTGCTGCAGCGACCAGCGCAGCAGCAGCACCCTGCACGAACACCCCG 159
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                                                                                                                                                                       CTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGGCAGCAGCACCTGCACGAACACCCG
ANACGTCTCCGAGAAGGGGGGGGGGGTCATGTACGCCCGGGAAGTAGGACCTCGTCCAGTC
                                                                                                                  CCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAA 219
                                                          CCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Possible reversed clone: similarity seq primer: -40UP from Gibco High quality sequence stop: 457.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tumor Gene Index Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov/image/html/iresources.shtml
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bonaldo, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tissue Procurement: Christopher A. Moskaluk, M.D., R. Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
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/clone="IMAGE:2949166"
/clone_lib="NCI_CGAP_GC6"
/tissue_type="pooled germ cell tumors"
/lab_host="DH10B"
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                                                                                                                                                                                                                 Bonaldo, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-GGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL, send email to:
                                                                                                                                                                                                                                                                                                                        (CGAP/BTGAP), Tumor Gene Index Unpublished (1998)
Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N. Louis,
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1 (bases 1 to 551)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index
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/clone-"IMAGE:3306972"
/clone_lb-"NCI_CGAP_Brn23"
/tlssue_type="911oblastoma (pooled)"
/lab_host-"DH10B"
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1 (bases 1 to 540)
1 (chases 1 to 540)
1 NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurola
Disorders and Stroke, Brain Tumor Genome Anatomy Project
(CGAP/BTGAP), Tumor Gene Index
Unpublished (1998)
Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N. Loui
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/clone="IMAGE:3440591"
/clone_lib="NCI_CGAP_Brn23"
/tissue_type="glioblastoma (pooled)"
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                AGGAGGAGAGACAGAGAAAAGAAAAACACAGCATGAGAAACACAGTAAATGAATAAAACC
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Plate: 06 row: h column:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="by06h03"
/clone_lib="Human Lens cDNA (Un-normalized, unamplified):
BY"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /dev_stage="Adult"
/lab_host="EMDH10B"
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                                                                                     CACCCGCCGAAACTGCTGCGAGGACCACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAG
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                                GTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGT
                                                                                                                                                            CAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGGCAGCAGCACAGCACCTGCACGAA 153
GTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGT
                                                                   CACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAG
                                                                                                                                        CAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGGCAGCACCAGCACCTGCACGAA 468
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                                                                                                                                                                                                              512;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrat Mammalla; Eutheria; Primates; Catarrhini; Hominid (bases 1 to 527)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy
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AW338938.1
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                                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ife Technologies catalog #: 11548-013
DNA Sequencing by: Washington University Genome Sequencing Center
                                                                                                                                                                                                                                                                                                  103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             quality sequence stop: 420
                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                             /note="Organ: pancreas; Vector: pCMV-SPORT6; Site_1: Sall;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.72 kb. Life Technologies catalog #:
                                                                                                                                                                                                                                                                                                                  11548-013"
                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="NCI_CGAP_Pan1"
/tissue_type="adenocarcinoma"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="IMAGE: 2878903"
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Pred. No. 0;
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                                                                                                                                                                                                                                                                                                                                                                                               found through the I.M.A.G.E. Consortium/LLNL www-bio.llnl.gov/bbrp/image/image.html Insert Length: 1161 Std Error: 0.00 Seq primer: -40UP from Gibco
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 515)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bonaldo, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R. Emmert-Buck, M.D., Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 quality sequence stop: 444.
Location/Qualifiers
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/note="Vector: PT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Plasmid DNA from the normalized library NCI_CGAP_GC4 was prepared, and scircles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified CDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1257096-1258631, 1469064-1470983, and 1475592-1476743).
                                                                                                                                                                                        /clone_lib="NCI_CGAP_GC6"
/tissue_type="pooled germ cell tumors"
/lab_host="DH10B"
                                                                                                                                                                                                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2009444"
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BASE COUNT ORIGIN

103

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97

Query Match Best Local S Matches 509

Similarity

73.4%;

Score 459; Pred. No. 0;

Conservative

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TTTGCCACAAGCAAGAGAATTTAACACTGTTTCAAACCCGGGGGAGTTGGCTGTGTTA
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1 (bases 1 to 690)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index (CGAP/BTGAP), Tumor Gene Index (Daybullashed (1998)
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Burt Feuerstein, M.D., Mark Israel, M.D.
cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing C
Clone distribution: NCI-CGAP clone distribution information
found through the I.M. A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                                                                                                                                   AW149665

Kf40h05.xl NCI_CGAP_Brn50 Homo sapiens cDNA clone IMAGE:2620569 3/
Similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
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                                                                                                                                                    AAACCCGGGGGAGTTGGCTGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG
                                                                                                                                                                                                                              GCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGAGGATCATGT
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                                                                                                                                                                                                                                                                                                    AACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTT
                                                                                                                                                                                                                                                                                                                                                        GAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACAC
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                                                                                                                                 AAACCCGGGGGAGTTGGCTGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG
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AW075598
xb24e07.x1
similar to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_"IMAGE:2620569"
/clone_lib="NCI_CGAP_Brn50"
/tissue_type="medulloblastoma"
/lab_host="DH10B (phage resistant)"
/lab_host="DH10B (phage resistant)"
/lab_host="DH10B (phage resistant)"
/note="Organ: brain; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from medulloblastoma tumor tissue, and was then primed with a Not I - ollgo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. This library is normalized. Library constructed by Bento Soares and M. Fattma Bonaldo. "
42 a 196 c 160 g 191 t l others
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/db_xref="taxon:9606"
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500 bp mRNA linear EST 13-OCT-1999
NCI_CGAP_Kidl3 Homo sapiens cDNA clone IMAGE:2577252 3'
SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
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99.58;
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RESULT 14 AW149665/c LOCUS

DEFINITION

wrksy.; mRNA sequence.

AW149665.1 GI:6197561

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35

AAGAAAGACCATTAAAATGCTTTAGACAGTG

606 σ

SOURCE ORGANISM

Homo sapiens

KEYWORDS VERSION ACCESSION

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REFERENCE

AUTHORS TITLE

COMMENT

Contact: Robert Strausberg, Ph.D.

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Best Local :
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                GCCATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGC
                                                                                                                                                                                                                                                                                                                                                   AGTAAATAAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAA 131
                                                                                       AGTAAATGAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCCTACTGGCCAGGAA 478
                                                                                                                                        CCTGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACAC
                                                                                                                                                                                                            GCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGC
                                                                                                                                                                                                                                                                              AGGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCA 298
                                                                                                                                                                                                                                                                                                                                                                                    ACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGG 238
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Genome Sequencing Center Clone distribution information can Clone distribution: NCI-CGAP clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tissue Procurement: Chris Moskaluk, M.D., Ph.D., Michael R. Emmert-Buck, M.D., Ph.D. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: Christa Prange, Th. I.M.A.G.E. Consortium DNA Sequencing by: Washington University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GPR39.;, mRNA sequence.
AW075598
AW075598.1 GI:6030596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 500)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note="Organ: kidney; Vector: pCMV-SPORT6; Site_1: Sal
Site_2: NotI; Cloned unidirectionally. Primer: Oligo
Library constructed by Life Technologies."
a 139 c 114 g 148 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="NCI_CGAP_Kid13" /tissue_type="2 pooled Wilms' tumors, one primary and metastatic to brain" /lab_host="DH10B"
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Pred. No. 0;
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TITLE
                                                                                                                                            Matches 425;
                                                                                                                                                                 Best
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GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
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Insert Length: 1462 Std Error: 0.00
Seq primer: -40UP from Gibco
High quality sequence stop: 444.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 591)

NCI/NIMDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
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                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                            109
                                                                                                                                          Conservative
                                                                                                                                                                                                                                      T 3']; double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized, and was constructed by Bento Soares and M.Fatima Bonaldo."

179 c 147 g 155 t 1 others
                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE:2431501"
/clone_lib="NCI_CGAP_Brn25"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                           /tissue_type="anaplastic oligodendroglioma"
/lab_host="DH10B"
                                                                                                                                                             68.0%;
100.0%;
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                                                                                                                                                             Score 425; E
Pred. No. 0;
                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                 cDNA Library Preparation: Ling Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov Plate: LLCM285 row: g column: 09
                                                                                                                                                                                                                                                                                                                                                                                                                           Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryola; Metazoa; Chordata; Craniata; Vertebrata; Mammalla; Eutherla; Primates; Catarrhini; Hominidae; 1 (bases 1 to 676)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collections (1988)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1999)
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                                          ECORI; CDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GCCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-CDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies)." a 172 c 199 g 137 t
                                                                                                                                                                  /tissue_type="melanotic melanoma"
/lab_host="DH10B (phage-resistant)"
/note="Organ: skin; vector: pOTB7; Site_1: xhoi; Site_2:
                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:3617696"
/clone_lib="NIH_MGC_20"
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IMAGE:3617696 5',
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                               source
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Score

Length

Unpublished (1998)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov

Tissue Procurement: David N. Louis,

M.D.,

Myrna R.

Rosenfeld

M.D.,

3

Fatima

cDNA Library Preparation: M. Bento Soares, Ph.D.,

CONA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome
Clone distribution: NCI-CGAP clone distribution
found through the I.M.A.G.E. Consortium/LLNL at:

Sequencing Center information can be

primer: -40m13 fwd. ET from Amersham quality sequence stop: 425

/organism="Homo sapiens" /db\_xref="taxon:9606"

1. .467

ocation/Qualifiers

Bonaldo, Ph.

Ph.D.

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutherla; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 467)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index

EST

human

AI150931.1 sequence.

GI:3679400

Homo sapiens

qb53c04.x1 similar to

467 bp mRNA linear EST 26-OCT-1998 NCI\_CGAP\_Brn23 Homo sapiens cDNA clone IMAGE:1703814 3' SW:NTR1\_RAT P20789 NEUROTENSIN RECEPTOR TYPE 1 ;, mRNA

AI150931

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TAAAT
                      TAAAT 425
                                            CAGCGACAGGCGGCAGCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC
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                                                                                                                   CATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCAGACACGAGCAGGAGGAGAGACAGAGAAAAAAAAACACAGCATGAGAACACAGTAA
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                                                                                                                                                                                                  UI-E-DWO-agm-j-01-0-UI.Sl UI-E-DWO Homo sapiens cDNA UI-E-DWO-agm-j-01-0-UI 3', mRNA sequence.
                                                                                   Eukaryota, Metazoa; Chordata, Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 538)
Bonaldo, M.F., Lennon, G. and Soares, M.B.
                                                                                                                                                                          EST
Contact: Soares, MB
Program for Rat Gene Discovery and Mapping
                                                            discovery
                                                                     Normalization and subtraction: two
                                                                                                                                            Homo sapiens
                                                                                                                                                                                    BM667957.1 GI:18975788
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                                            Res. 6 (9), 791-806 (1996)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE:1703814"
/clone_lib="NCI_CGAP_Brn23"
/tissue_type="glioblastoma (pooled)"
/lab_host="DH10B"
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AGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAGAGAGAATTTAACACTGTTTCAAA
                                           CATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTC
                                                                CATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTC 494
                                                                                                                                                           GGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACG
                                                                                                                                                                                                                                                                                                                                                                                                                                               GGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACG
                                                                                                                                   TCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAG
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Tissue Procurement: Dr. Gregg Hageman
CDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iowa
CDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa
DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
Clone Distribution: Researchers may obtain clones from Research
Genetics (www.resgen.com).
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451 Eckstein Medical Research Building Iowa City,
Tel: 319 335 8250
Fax: 319 335 9565
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/note="Organ: eye; Vector: pT7T3-Pac (Pharmacia) with a
modified poolylinker; Site_1: EcoR I; Site_2: Not I;
UI-E-DW0 is a cDNA library containing the following
tissue(s): lens. The library was constructed according to
Bonaldo, Lennon and Soares, Genome Research, 6:791-806,
1996. First strand cDNA synthesis was primed with an
oligo-dT primer containing a Not I site. Double stranded
cDNA was ligated to an EcoR I adaptor, digested with Not
I, and cloned directionally into pT7T3-Pac vector. The
oligonucleotide used to prime the synthesis of
first-strand cDNA contains a library tag sequence that is
located between the Not I site and the (dT).B tail. The
sequence tag for this library is CGATTAGCGA. This library
was created for the program, Gene Discovery in the Visual
System, supported by National Eye Institute (NEI).
TAG TISSUFF-human lens
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TAG_SEQ=CGATTAGCGA"
144 c 122 g
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/clone_lib="UI-E-DW0"
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/db_xref="taxon:9606"
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RESULT 20
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                                                                                                                                                                                84
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                                CCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGT 263
                                                                                          CCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGA 203
                                                                                                                                          CATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGGCAGCAGCA 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCCGGGGGAGTTGGCTGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG 606
CCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGT
                                                                       CCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGA 418
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
cDNA Library Preparation: M. Bento Soares, Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 537)
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Insert Length: 1394 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 452.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1997)
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                     and Eco RI sites of the modified pT7T3 vector. Libr went through one round of normalization to Cot5, an constructed by Bento Soares and M. Fatima Bonaldo. 147 c 126 g 159 t
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/clone_lib="Soares_testis_NHT"
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/db_xref="taxon:9606"
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similar to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: L. Jeffrey Medeiros, M.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                          High quality sequence stop: 395.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                www-bio.llnl.gov/bbrp/image/image.html
Seq primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                   DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EST
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                                    56
                              /clone_lib="NCI_CGAP_AA1"
                                                                                                                                                                                                                 /db_xref="taxon:9606"
/clone="IMAGE:1113710"
                                                                                                                                                                                                                                                         /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           396 bp mRNA linear EST 06-OCT-1997 NGI_CGAP_AA1 Homo sapiens cDNA clone IMAGE:1113710 3' SW:NTR_RAT P20789 NEUROTENSIN RECEPTOR ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Inc.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 David B. Krizman,
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                                                                                                                                                                                                                                                                                                                               (CGAP/BTGAP), Tumor Gene Index
Unpublished (1998)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld
                                                                                                                                                               found through the i.m. for that www-bio.llnl.gov/bbrp/image/image.html rnsert Length: 1183 Steror: 0.00
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1 (bases 1 to 452)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       452 bp mRNA linear EST 17-MAR-1999 qy93f04.x1 NCI_CGAP_Brn25 Homo sapiens cDNA clone IMAGE:2019583 3'similar to SW:NTR1_RAT P20789 NEUROTENSIN RECEPTOR TYPE 1 ;, mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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DNA Sequencing by: Washington University Genome
Clone distribution: NCI-CGAP clone distribution
found through the I.M.A.G.E. Consortium/LLNL at:
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                                                                                                                                         quality sequence stop: 451.
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/clone="IMAGE:2019583"
/clone_lib="NCI_CGAP_Brn25"
/tissue_type="anaplastic ol
                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                      ocation/Qualifiers
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                                                                                                                                                                                                                                                                     sequence.
               Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
Emmert-Buck, M.D., Ph.D.
                                                                                Tumor Gene Index
Unpublished (1997)
                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 396)
                                                                                                                                                                                                                                                                                               AI499630 396 bp mRNA linear EST 12-MAY-1999 to03b07.xl NCI_CGAP_Ut2 Homo sapiens cDNA clone IMAGE:2177941 3' similar to SW:NTRL_RAT P20789 NEUROTENSIN RECEPTOR TYPE 1 ;, mRNA
                                                                  Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                    Homo sapiens
                                                                                                                  NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Ana
cDNA Library Preparation: Life Technologies, Inc
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Pred. No. 0;
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                                                                                                                Genome Anatomy Project (CGAP),
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DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 534) NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
                                                                  Homo sapiens
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                                                                                                                                                  ;, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="NCI_CGAP_Ut2"
/tissue_type="moderately-differentiated endometrial adenocarcinoma, 3 pooled tumors"
/lab_host="DH10B"
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/db_xref="taxon:9606"
/clone="IMAGE:2177941"
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                                                                                  AI765236 456 bp mRNA wi72h08.xl NCI_CGAP_Kidl2 Homo sapiens cl similar to SW:GP39_HUMAN 043194 PUTATIVE
                                  GPR39. ;, mRNA sequence.
AI765236
AI765236.1 GI:5231745
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밁 δ Вb Ş ORIGIN

58

BASE COUNT

FEATURES

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358 156 298 216 238 276 178 336 118 396

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36

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RESULT 24

VERSION

p mRNA linear EST 21-DEC-1999 sapiens cDNA clone IMAGE:2398911 3'4 PUTATIVE G PROTEIN-COUPLED RECEPTOR

ACCESSION

REFERENCE AUTHORS

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Query Match
Best Local Similarity
                  361 TGGGCAGACACGAGCAGGAGGGGAGAGACAG 390
                                                                           121 CAGCGACAGGGCGCACCACACCTGCACCGCAACACCCGCCGAAACTGCTGCGAGGACAC
                                                                                                                                                                                                                                                                                                                                                                                                  330 GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                390 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
                                                                                                                                                         150 GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC
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TGGGCAGACACGAGCAGGAGGGGAGACAG
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Clone distribution: NCI-CGAP clone distribution information can found through the I.M.A.G.E. Consortium/LLNL, send email to:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       High quality sequence stop: 'Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tissue Procurement: Chris Moskaluk, M.D., Ph.D., Michael R. Emmert-Buck, M.D., Ph.D. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: Christa Prange, Th. I.M.A.G.E. Consortium DNA Sequencing by: Washington University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         National Cancer Institute,
Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 info@image.llnl.gov
Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="Organ: kidney; Vector: pCMV-SPORT6; Site_
Site_2: NotI; Cloned unidirectionally. Primer:
Library constructed by Life Technologies."
a 172 c 143 g 121 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE:3146086"
/clone_lib="NCI_CGAP_Kid13"
/tissue_type="2 pooled Wilms' tumors, one primary and one
metastatic to brain"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                           GTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTTGGGTTTGGGCATCCAGCATAC 336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTA 216
TTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGTTTGGACTTGACAGCTTCT 516
                                                                                                                                                                                                 GAAAAACACAGCATGAGAAACACAGTAAAATGAATAAAAACCATAAAAATATTTTAGCCCCTCTG 456
                                                                                                                                                                                                                                                                                     GAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCA 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCA 276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTA 397
                                           TTTGCCACAAGCAAGAGAATTTAACACTGTTTCAAACCCGGGGGAGTTGGCTGTTTA 576
                                                                                                                                                                            GAAAAACACAGCATGAGAACACAGTAAATAAATAAAACCATAAAATATTTAGCCCCTCTG 157
                                                                                                                                                                                                                                                                GTCGTGCTTGGGTTTGGCCGAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATAC
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1 (Dases 1 to 456)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 www-bio.llni.gov/bbrp/image/image.html
Insert Length: 961 Std Error: 0.00
Seq primer: -40UP from Gibco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tumor Gene Index
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 quality sequence stop: 453.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note-"organ: kidney; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Plasmid DNA from the normalized library NCI_CGAP_Kid5 was prepared, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1323912-1325831, 1471568-1472903 and 1492104-1493255). "Subtraction by Bento Soares and M. Fatima Bonaldo."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone_lib="NCI_CGAP_Kid12"
/tissue_type="2 pooled tumors (clear cell type)"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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99.8%;
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Pred. No. 0;
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VERSION
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Best Local S
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                                                                                                                                                                                                                   237 GGAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCG 296
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ACAGTAAATGAATAAAACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGG 476
                                   CAGCCATGATCCTCCGAATCTGGTTTGGGCATCCAGCATACGGCCAATGTCACAACAATCA 120
                                                                                                                                                           CAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCA 356
                                                                                                                                                                                                GGAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCG 60
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                                                                           GCCCTGGGCAGACACGAGCAGGAGGGGAGAGACAGAGAAAAGAAAAACACAGCATGAGAAC 416
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NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurological
Disorders and Stroke, Brain Tumor Genome Anatomy Project
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381 bp mRNA linear EST 16-OCT-2000 7171h01.yl NCI_CGAP_Brn20 Homo sapiens cDNA clone IMAGE:3340177 5' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Putative full length read
The vector to vector length is 382
Seq primer: -40RP from Gibco.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CDNA Library Preparation: David B. Krizman, Ph.D. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL, send email to: info@image.llnl.gov
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Other_ESTs: 7i71h01.x1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tissue Procurement: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                                                                                                                                                                  /note="Organ: brain; Vector: pAMP1; mRNA made frolligodendroglioma tissue, cDNA made by oligo-dT Directionally cloned. Size-selected on agarose average insert size 500 bp. Primary library, non-amplified."

81 c 94 g 83 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone_lib="NCI_CGAP_Brn20"
/tissue_type="oligodendroglioma"
/dev_stage="adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone="IMAGE: 3340177"
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                                                                                                                                                                                                                                                                                                     Score 369;
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                                                           238 GAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGC 297
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Robert Jenkins, M.D., Mark Israel,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleosto; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 371)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BF054680 mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BF054680 371 bp mRNA linear EST 16-OCT-2000 7169f08.yl NCI_CGAP_Brn20 Homo sapiens cDNA clone IMAGE:3339975 5' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High quality sequence stop: 361. Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (1998)
Other_ESTs: 7169f08.x1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      info@image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         found through the I.M.A.G.E. Consortium/LLNL, send email
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CDNA Library Preparation: David B. Krizman, Ph.D.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
                                                                                                                      Similarity
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                                                                                                                                                                                                                              average insert size 500 bp. non-amplified. "
                                                                                                                                                                                                                                           /note-"Organ: brain; Vector: pAMP1; mRNA made fx oligodendroglioma tissue, cDNA made by oligo-dT Directionally cloned. Size-selected on agarose average insert size 500 bp. Primary library,
                                                                                                                                                                                                       non-amplified.
                                                                                                                                                                                                                                                                                                                           /lab_host="DH10B"
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/tissue_type="oligodendroglioma"
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/clone="IMAGE:3339975"
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W938h03:x1 Soares_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone
IMAGE:2367413 3' similar to SW:GP39_HUMAN 043194 PUTATIVE G
PROTEIN-COUPLED RECEPTOR GPR39. ; mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov This clone is available royalty-free through LLNL ; contact the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 578)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Insert Length: 806 Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                quality sequence stop: .
Location/Qualifiers
              Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 5 libraries. The pools consisted of the following libraries and cloneIDs: Soares NbHFP pool 1: 309384-310919, 323208-325895 Soares NbHPP pool 1: 145032-147335, 147720-148103, 148872-149255, 15002-150407, 151176-152327 Soares NbHFP apol 1: 758280-760583, 772104-774407 Soares NbHPA pool 1: 304776-306311, 320136-323823, 326280-236663 Soares NbHOT pool 1: 723720-726407, 739080-740999 Subtraction by Bento
                                                                                                                                                                                                                                           /note-"Organ: pooled; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from five normalized libraries were mixed, and ss circles were made in vitro.
                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:9606"
/clone="IMAGE:257413"
/clone_lib="Soares_NSF_F8_9W_OT_PA_P_S1"
/lab_host="DH10B"
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mes 415; Conserv
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                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
Emmail: cgapbs-r@mail.nih.gov
Inssue Procurement: Chris Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D. cDNA Library Preparation: Life
Technologies, Inc. cDNA Library Arrayed by: Christa Prange,
I.M.A.G.E. Consortium DNA Sequencing by: Washington Universi
                                                                       Seq
                                                                                                                            Genome Sequencing Center Clone distribution Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  AW087372

432 bp mRNA linear EST 15-OCT-1999 xb19902.xl NCI_CGAP_Kidl3 Homo sapiens cDNA clone IMAGE:2576786 3' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 432)
                                                                                                                                                                                                                                                                                                                                                                                                                          GPR39.;, mRNA sequence.
                                                                                     Possible reversed clone: similarity
                                                                                                                www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                               Unpublished (1997)
                                                                                                                                                                                                                                                                                          NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                Tumor Gene Index
                                                                       primer: -40UP from Gibco
                                          quality sequence stop: 306
Location/Qualifiers
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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TITLE
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320 TTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAGGA 379
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nes 347; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGT
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                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                           CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC Clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov Plate: LLCM855 row: o column: 21 High quality sequence stop: 723.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4026092"
/clone_lib="WIH_MGC_20"
/tissue_type="melanotic melanoma"
/lab_host="DHIOB (phage-resistant)"
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Site_2: Notl; Cloned unidirectionally. Primer: Oligo dT
Library constructéd by Life Technologies."
1 16 c 89 g 134 t 1 others
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/tissue_type="2 pooled Wilms' tumors, one primary
metastatic to brain"
/lab_host="DH10B"
                                                                                                                                                                               Location/Qualifiers
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RESULT 31
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BF125134
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                                                                                                                           High quality sequence stop: 711.
Location/Qualifiers
                                                                                                                                                                       Plate: LLCM853 row: p column: 13
                                                                                                                                                                                        cDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.
                                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                     NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="InhAGE:4025340"
/clone_lib="NIH_MGC_20"
/tissue_type="melanotic melanoma"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GI:10964174
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Pred. No.
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RESULT 32
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Best Local
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                      This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Possible reversed clone: similarity on wrong strand Possible reversed clone: polyT not found Seg primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 387)

Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost, S., Krizman, D., Kucaba, T., Lacy, M., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AI272281 387 bp mRNA linear EST 18-NOV-1998 ap23c03.x1 Schiller oligodendroglioma Homo sapiens cDNA clone IMAGE:1956196 3' similar to SW:NTR1_RAT P20789 NEUROTENSIN RECEPTOR TYPE 1 ;contains TAR1.t3 TAR1 repetitive element ;, mRNA sequence.
                                                                                                                                          Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                  Washington University Scho
4444 Forest Park Parkway,
                                                                                                                          Email: est@watson.wustl.edu
                                                                                                                                                                                                                           Contact: Wilson RK
                                                                                                                                                                                                                                                               White,Y., Wylie,T., Waterston,R. WashU-NCI human EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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    High quality
                                                                                                                                                                                                                                            Unpublished (1997
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/note="Organ: skin; Vector: pOTB7; Site_1: XhoI; Site_2:
ECORI; CDNA made by oligo-dT priming. Directionally
cloned into EcoRI/XhoI sites using the following 5'
adaptor: GGCACGAG(G). Size-selected >500bp for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubln (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
a 192 c 255 g 171 t
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sequence stop: 386
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Pred. No.
                                                                                                                                                                                    School of Medicine way, Box 8501, St. 1
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RESULT 33
AW191974/c
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                                                                                                                                                                                                                                                                                                                                                                                                                GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
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                                                                                                                                                                                                                                                                                   GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC 300
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                                                                                                                                                                                                                                                                                                                                                                                                      GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                                                    AW191974
x178a02.x1 NCI_CGAP_Pan1
similar to SW:GP39_HUMAN
Unpublished (1997)
Contact: Robert Strausberg,
                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 454)
                                                                                                            EST
                                                                                                                    GPR39. ;, mRNA sequence.
AW191974
AW191974.1 GI:6470673
                       Tumor Gene Index
                                                                                      Homo sapiens
                                 National Cancer Institute,
                                        NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
                                                                                                  human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="oligodendroglioma"
/dev_stage="44 years"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="Schiller oligodendroglioma"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /lab_host="SOLR"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 338;
Pred. No.
                                                                                                                                                   454 bp mRNA linear EST 29-NOV-1999
Homo sapiens cDNA clone IMAGE:2680778 3'
043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                 Cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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                                 Anatomy
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                               Project (CGAP),
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173 GAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAG 232
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  A1423162 480 bp
tf32c12.x1 NCI_CGAP_Brn23 Homo s
similar to SW:NTR1_RAT P20789 NE
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DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       High quality sequence stop: 395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      www-bio.llnl.gov/bbrp/image/image.html
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             reversed clone: similarity on wareversed clone: polyT not found
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/db_xref="taxon:9606"
/clone="IMAGE:2680778"
/clone_lib="NCI_CGAP_Pan1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /tissue_type="adenocarcinoma"
/lab_host="DH10B"
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p mRNA linear EST 30-MAR-1999 sapiens cDNA clone IMAGE:2097910 3'
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REFERENCE

ORGANISM

AUTHORS TITLE

JOURNAL

Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov

Tissue Procurement: David

Bonaldo,

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133 GCAGCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAG 192
                      TCAGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAGAGAGAATTTAACACTGTTTCA
                                                                                  ACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTT 492
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                                                              ACCATAAAATATTTAGCCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTT
                                                                                                                                                   Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 480)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
NATIONAL Cancer Institute / National Institute of Neurola Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index Unpublished (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cDNA Library Arrayed by: Greg Lennon, Ph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cDNA Library Preparation: M. Bento Soares, Ph.D.,
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the Not I and Eco RI sites of the modified
Library is normalized, and was constructed
Soares and M.Fatima Bonaldo."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /tissue_type="glioblastoma (pooled)"
/lab_host="DH10B"
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                    GAACACAGTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGC 472
GAACACAGTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGC
                                                                                    ATCAGCCCTGGGCAGACACGAGCAGGAGGGGAGAGACAGAAAAGAAAAACACAGCATGA 195
                                                                                                                             ATCAGCCCTGGGCAGACGAGCAGGAGGGGAGAGAGAAAAGGAAAAACACAGCATGA 412
                                                                                                                                                                       GCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACA 255
                                                                                                                                                                                                                                                            314;
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AA665640
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This clone is available royalty-free through LLNL; MAGE Consortium (info@image.llnl.gov) for further i Seq primer: -40ml3 fwd. Er from Amersham High quality sequence stop: 325.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         High quality sequence stop:
Location/Qualifiers
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Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1997)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                     78
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/sex="pooled (6)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone="IMAGE:1126871"
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/db_xref="taxon:9606"
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BASE COUNT

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Local Similarity

50.4%;

Score 315; Pred. No. 0; Mismatches

Conservative

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Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost, S., Krizman, D., Kucaba, T., Lacy, M., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B., White, Y., Wylie, T., Waterston, R. and Wilson, R. Washi, NCI human EST Project
/note="vector: pSPORT1; Site_1: Sall; Site_2: Not1; RNA
was prepared from a pool of 6 anonymous wilms' tumor RNAs.
RNA was prepared by acid-phenol, followed by one round of
oligo dT selection. cDNA library preparation was with
the BRL/Life Tech. Superscript Plasmid System. An
oligo-dT NotI primer for first strand systhesis generated
geggegecc(t)n at the 3' end of the clones. A 5' Sall
adaptor was used with sequence 5'-gtcgaccacgcgtccg-3'.
Resulting cDNAs were size selected (average size 2 kb),
NotI digested, and ligated into NotI/SalI-cut pSPORT1.
Library was constructed by Dr. Manfred Gessler."
78 a 81 c 74 g 104 t
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FEATURES

ity sequence stop: 372. Location/Qualifiers

source

/clone="IMAGE: 2097910" /clone\_lib="NCI\_CGAP\_Brn23"

/db\_xref="taxon:9606" /organism≃"Homo sapiens"

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                GAACACAGTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGC 472
                                                                                        GAACACAGTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGC 136
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1 (bases 1 to 354)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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AA970361.1 GI:3145874
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                               /note-"Organ: pooled; Vector: pT773D-Pac (Pharmacia) with a modified polyinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from three normalized libraries (fetal lung NbHL19W, testis NHT, and B-cell NCI_CGAP_GCB1) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-302087, 682632-687239, 726408-728711, and 729096-731399. Subtraction by Bento Soares and M. Fatima Bonaldo. "
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="Soares_NFL_T_GBC_S1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone="IMAGE:1580506"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
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314 286 1810
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Email: est@watson.wustl.edu
WARNING: There is evidence that suggests that the 384-well parent
plate of this clone contains both human and mouse derived clones.
Thus, the origin of this clone is uncertain. This caution should kept in mind should you use this clone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 373)
Hillier, L., Lennon, G., DuBuque, T., Favello, A., Gish, W., Hawkins Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Le, N., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M. Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Insert Length: 427 Std Error: 0.00
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                                                                   /organism="Homo sapiens"
/db_xref="GDB:3923283"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                           /sex="female"
                                                                                                                                                                                                                                                                                                                           /clone_lib="Stratagene ovarian cancer (#937219)"
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Score 314;
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Znl7f09.S1 Stratagene neuroepithelium NT2RAMI 937234 Homo sapiens
cDNA clone IMAGE:547721 3', mRNA sequence.
AA084249
AA084249.1 GI:1626369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hillier, L., Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Le, M., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M. Generation and analysis of 280,000 human expressed sequence tags Genome Res. 6 (9), 807-828 (1996)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
/clone_lib="Stratagene neuroepithelium NT2RAMI 937234"
/dev_stage="Ntera-2/RA+MI neuroepithelial cells"
/lab_host="SOLR (kanamycin resistant)"
/note="Vector: pBluescript SK-, Site_1: EcoRI; Site_2: XhoI; Cloned unidirectionally. Primer: Oligo dT. NT2
(Ntera-2/c1.D1) precursor cells induced with Retinoic Acid for 1 week, followed by 3 weeks in mitotic inhibit
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Tel: 301 402 3452
Fax: 301 496 0078
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BF726644
                                                                                                                                                                                                                                                                                           Email: graeme@helix.nih.gov
Plate: 09 row: g column: 04
Seg primer: M13RP1 reverse primer (ABI).
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                                                                                                                                                                                                                                                                                                                                                                                                                            Section on Molecular Structure and Function National Eye Institute
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                                          /clone_lib="Human
BY"
  /note="Organ: Eye; Vector: pCMVSPORT6; Two human lenses from different adults (both approximately 40 years old) together yielded 20ug of total RNA and 150ng mRNA for c
                                                                                                                                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                     /tissue_type="Lens"
/dev_stage="Adult"
/lab_host="EMDH10B"
                                                                                                                                                                                 /clone="by09g04"
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FEATURES

Location/Qualifiers /organism="Homo sapiens"

source

/db\_xref="GDB:3926237" /db\_xref="taxon:9606" /clone="IMAGE:547721"

COMMENT

Contact: Wilson RK
Washington University Sch
4444 Forest Park Parkway,

School of Medicine

Box 8501,

St. Louis,

Tel: 314 286 1800 Fax: 314 286 1810 Email: est@watson.wustl.edu

TITLE JOURNAL MEDLINE

Genome Res. 97044478

REFERENCE

(bases 1 to 412)

AUTHORS

SOURCE VERSION ACCESSION KEYWORDS

ORGANISM

Homo sapiens numan

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(Replate #2). Average insert size: 1.1 kb; Uni-ZAP Vector; ~5' adaptor sequence: 5' GAATTCGGCACGAG 3' adaptor sequence: 5' CTCGAGTTTTTTTTTTTTTTTT 3'" 110 c 87 g 126 t
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1 (bases 1 to 349)

Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost, S., Kizman, D., Kucaba, T., Lacy, M., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B., White, Y., Wylle, T., Waterston, R. and Wilson, R.

Washu-NCI human EST Project
                                              Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL;
IMAGE Consortium (info@image.llnl.gov) for further i
Insert Length: 792 Std Error: 0.00
                                                                                                                                                                                                  Washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis,
                                                                                                                                                                                                                                                 Unpublished (1997)
Contact: Wilson RK
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zp04h05.rl Stratagene ovarian clone IMAGE:595449 5', mRNA se
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408 bp mRNA linear EST 25-J
7f75g10.x1 Lupski_dorsal_root_ganglion Homo sapiens cDNA clc
IMAGE:3302971 3' similar to SW:Gp39_HUMAN 043194 PUTATIVE G
PROTEIN-COUPLED RECEPTOR GPR39. ; mRNA sequence.
BG057775
                                                                                                                                                                                                                                                                 EST
                                                                         Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Dr. James Lupski
                                                                                                                       Unpublished (1997)
                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                 Homo sapiens
                                                                                                       Contact: Robert Strausberg, Ph.D.
                                                                                                                                                      National Cancer Institute, Cancer
                                                                                                                                                                    NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
                                                                                                                                                                                                                                                                              BG057775.1 GI:12523767
found through the I.M.A.G.E. Consortium/LLNL
           CDNA Library Preparation: Lupski Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Washington University Genome Sequencin
Clone distribution: NCI-CGAP clone distribution informati
                                                                                                                                      Tumor Gene Index
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
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/sex="female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /dev_stage="adult, 64 years"
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100.08;
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                              Sequencing Center
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FEATURES

Seq p

info@image.llnl

primage.lini.gov primer: -40UP from Gibco quality sequence stop: Location/Qualifiers

source

/sex="male"

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GTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA 479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAAACACACA 419
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NCI/NINDS-GGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of N

Disorders and Stroke, Brain Tumor Genome Anatomy Pr
                                                                                                                                                                                                                   AI566797
420 bp mRNA linear EST 12-MAY-1999 tn23g08.xl NCI_CGAP_Brn25 Homo sapiens cDNA clone IMAGE:2168510 3' similar to SW:NTR1_RAT P20789 NEUROTENSIN RECEPTOR TYPE 1 ;, mRNA
                                                                                   Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5'-GACTAGTTCTAGATCGCGAGCGGCCCT(15)-3'. Size selected 1 kb for average insert length 1.7 kb. This is a primary library, non-amplified Library constructed by Life Technologies and donated by J. Lupski, M.D./Ph.D. (Baylc College of Medicine) and is available through Life Technologies."

79 g 128 t 3 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              //note-"Vector: pCMV-SPORT6 (Life Technologies); Site_1:
Not1; Site_2: Sal1; cDNA made by oligo-dT priming.
Directionally cloned using the following adaptors:
5'-TCGACCCACGCGTCCG-3' and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:3302971"
/clone_lib="Lupski_dorsal_root_ganglion"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /dev_stage="adult, 36
/lab_host="DH10B"
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/dev_stage="adult, 36 yr"
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Pred. No. (
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                                                                                                                                                                                                                                                                                                   AAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGT 497
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                                                qy56f02.x1
similar to
sequence.
AI363261
AI363261.1
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N. Louis, M.D., Myrna R.
Ph.D.
                                                                               AI363261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 1148 Std Error: 0.00
Seq primer: -40UP from Gibco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: washington University Genome Clone distribution: NCI-CGAP clone distribution found through the I.M.A.G.E. Consortium/LLNL at:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="IMAGE:2168510"
/clone_lib="NCI_GAP_Brn25"
/tissue_type="anaplastic oligodendroglioma"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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                                                NCI_CGAP_Brn23 Homo
SW:NTR1_RAT P20789 N
GI:4114882
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99.78;
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Pred. No. 0;
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                                                                                 475 bp
                                              bp mRNA linear EST 16-FEB-1999
o sapiens cDNA clone IMAGE: 2016027 3'
NEUROTENSIN RECEPTOR TYPE 1 ;, mRNA
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5

Query Match Best Local

Local Similarity nes 307; Conser

Conservative

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COUNT

79 g

REFERENCE AUTHORS TITLE

SOURCE ORGANISM VERSION KEYWORDS

Homo sapiens

ACCESSION

AI566797.1 EST. sequence. AI566797 RESULT 42 AI566797/c LOCUS

DEFINITION

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                                                                                                                                                                     GTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGC
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                                 TGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGTTGGACTTGACAGCTTCTTTT 519
                                                                 AAACCCAGCATGAGAACCCAGTAAATAAATAAAACCATAAAATATTTAGCCCCTCTGTTC
                                                                                                                                   CAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAGGAGGAGACAGAGAAAAGAA
                                                                                                                                                                                                       GTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGC
                                                                                                                                                                                                                                                                                                                                                                                                                  444;
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1 (bases 1 to 475)

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 814 Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seq primer: -40UP from Gibco
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/clone="IMAGE:2016027"
/clone_lib="NCI_GGAP_Brn23"
/tlssue_type="glioblastoma (pooled)"
/lab_host="DH10B"
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Pred. No. 0;
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CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG
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                                                                 CAGCGACAGGCGGCAGCACCAGCACCTGCACGAACCCCGCCGAAACTGCTGCGAGGACAC
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Plate: L/LR0388 row: o column: 21
High quality sequence stop: 665.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1999)
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NIH-MGC http://mgc.nci.nih.gov/
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AGENCOURT_8344092 NIH_MGC_110 Homo sapiens cDNA clone IMAGE:6248684
5', mRNA sequence.
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DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC clone distribution information can be
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/db_xref="taxon:9606"
/clone=lib="NIH_MCC_110"
/clone_lib="NIH_MCC_110"
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/tissue_type="ductal carcinoma, cell line"
/tissue_type="ductal carcinoma, cell line"
/note="Organ: pancreas; vector: pOTB7; Site_1: XhoI;
/note="Organ: pancreas; vector: pOTB7; Site_1: XhoI;
/site_2: ECORI; cDNA made by oligo-dT priming.
Directionally cloned into ECORI/XhoI sites using the
following 5 adaptor: GGCACCAG(G). Library constructed by
Ling Hong in the laboratory of Gerald M. Rubin (University
of California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies).
Note: this is a NIH_MCC_Library."
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                                      ACAGCATGAGAACACAGTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTG 463
                                                                                                                                                                                                                         TTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCCAAT 343
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Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
Emmert-Buck, M.D., Ph.D.
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/clone_llb="NCI_CGAP_Ov23"
/tlssue_type="tumor, 5 pooled (see description)"
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/db_xref="taxon:9606"
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GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compugen Ltd

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Minimum DB seq length: 0
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3: /cgn2_6/ptodata/2/pubpna/U
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. /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
. /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
. /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
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. /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

|                   |                    |                   |                    |                   |                   |                    |                    |                    |                    |                    |                   |                    |                    |                   |                   |                   |                     |                   | מל             |
|-------------------|--------------------|-------------------|--------------------|-------------------|-------------------|--------------------|--------------------|--------------------|--------------------|--------------------|-------------------|--------------------|--------------------|-------------------|-------------------|-------------------|---------------------|-------------------|----------------|
| c 19              | c 18               | c 17              | c 16               | c 15              | C 14              | c 13               | 12                 | c 11               | 10                 | с<br>9             | ი<br>8            | c 7                | 6                  | G                 | 4                 | ω                 | 2                   | 1                 | Result         |
| 38.6              | 39                 | 42.2              | 43.8               | 47.8              | 56.4              | 171.2              | 207.8              | 310.8              | 353                | 361                | 361.6             | 395.4              | 433.2              | 613               | 613               | 624               | 624                 | 624               | Score          |
| 6.2               | 6.2                | 6.8               | 7.0                | 7.6               | 9.0               | 27.4               | 33.2               | 49.7               | 56.5               | 57.8               | 57.9              | 63.3               | 69.3               | 98.1              | 98.1              | 99.8              | 99.8                | 99.8              | Query<br>Match |
| 1032              | 425                | 1137              | 894                | 2036              | 1464              | 201                | 381                | 349                | 390                | 373                | 1362              | 409                | 558                | 1619              | 1619              | 1897              | 625                 | 625               | Length         |
| 9                 | 10                 | 12                | 10                 | 10                | 10                | 10                 | 10                 | 10                 | 10                 | 10                 | 10                | 10                 | 10                 | 10                | 10                | 10                | 10                  | 10                | BB             |
| US-09-971-228-2   | US-09-960-352-1711 | US-10-037-616-3   | US-09-815-242-4029 | US-09-866-582-17  | US-09-149-045-1   | US-09-867-701-8894 | US-09-867-701-7826 | US-09-867-701-2409 | US-09-867-701-1532 | US-09-867-701-1516 | US-09-825-294-208 | US-09-867-701-4240 | US-09-867-701-2375 | US-09-825-294-211 | US-09-825-294-205 | US-09-825-294-214 | US-09-867-701-10876 | US-09-825-294-210 | ID             |
| Sequence 2, Appli | Sequence 1711, Ap  | Sequence 3, Appli | Sequence 4029, Ap  | Sequence 17, Appl | Sequence 1, Appli | Sequence 8894, Ap  | Sequence 7826, Ap  | Sequence 2409, Ap  |                    | Sequence 1516, Ap  | Sequence 208, App | Sequence 4240, Ap  | Sequence 2375, Ap  | Sequence 211, App | Sequence 205, App | Sequence 214, App | Sequence 10876, A   | Sequence 210, App | Description    |

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| 34.6                                   | 34.6                | ى س<br>4 4<br>6 6 | 34.8              |                    | 35.4               |                    |                    |                    | 35.8              | 36                | 36                | 36.4              | 37.2              | 37.2              | 38.2               | 38.4              | 38.6               | 38.6               | 38.6               | 38.6               | 38.6              | 38.6              | 38.6              |
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| 1200<br>372                            | 457                 | 1788<br>371       | 624               | 2462               | 2462               | 21860              | 3977               | 3977               | 2785              | 888               | 837               | 2314              | 3511              | 1743              | 332                | 1089              | 32207              | 32177              | 4063               | 3381               | 2776              | 1376              | 1149              |
| 10<br>12                               | 10                  | 10                | 10                | 10                 | 10                 | 10                 | 10                 | 10                 | 10                | 10                | 10                | 10                | 10                | 10                | 10                 | 10                | 10                 | 10                 | 10                 | 10                 | 12                | 9                 | 9                 |
| US-09-815-242-4132<br>US-10-052-545-11 | US-09-960-352-12765 | US-09-148-012-1   | US-09-804-551B-41 | US-09-880-107-2251 | US-09-964-824A-289 | US-09-764-877-3803 | US-09-764-869-2001 | US-09-764-869-1999 | US-09-925-302-122 | US-09-888-615-26  | US-09-885-441-1   | US-09-962-832-105 | US-09-753-831-3   | US-09-925-301-541 | US-09-867-701-9973 | US-09-993-844-14  | US-09-764-877-3250 | US-09-764-877-3251 | US-09-764-847-1091 | US-09-764-847-1092 | US-10-037-616-1   | US-09-971-228-4   | US-09-971-228-1   |
| Sequence 4132, Ap<br>Sequence 11, Appl | 12                  | Sequence 1, Appli | Sequence 41, Appl | Sequence 2251, Ap  | Sequence 289, App  | Sequence 3803, Ap  | Sequence 2001, Ap  | Sequence 1999, Ap  | Sequence 122, App | Sequence 26, Appl | Sequence 1, Appli | Sequence 105, App | Sequence 3, Appli | Sequence 541, App | Sequence 9973, Ap  | Sequence 14, Appl | Sequence 3250, Ap  | Sequence 3251, Ap  | Sequence 1091, Ap  | Sequence 1092, Ap  | Sequence 1, Appl1 | Sequence 4, Appli | Sequence 1, Appli |

## ALIGNMENTS

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; FEATURE:

NAME/KEY: misc_feature

; LOCATION: (1)...(625)

; OTHER INFORMATION: n = A,T,C or

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                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 210
LENGTH: 625
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 625; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN
FILE REFERENCE: 210121.484C5
                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
121 CAGCGACAGGCGGCAGCACCAGCACCTGCACGAACCCCGCCGAAACTGCTGCGAGGACAC 180
                   121 CAGCGACAGGCGGCAGCACCTGCACGACACCCGCCGAAACTGCTGCGGAGGACAC 180
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Sequence 10976, Application US/09867701

Retent No. US20020132237A1

GENERAL INFORMATION:

APPLICANT: A91ate, Paul A.

APPLICANT: Jones, Robert

APPLICANT: Jones, Robert

APPLICANT: Harlocker, Susan L.

APPLICANT: HORNATION: AND DIAGNOSTIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSTS OF OVARIAN CANCER

TITLE OF INVENTION: AND DIAGNOSTS OF OVARIAN CANCER

FILE REFERENCE: 210121.497

CURRENT APPLICATION NUMBER: US/09/867,701

CURRENT FILING DATE: 2001-05-29

NUMBER OF SEQ ID NOS: 10912

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 10876
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; Sequence 214, Application US/09825294

Patent NO. US20020004491A1

; GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.

; APPLICANT: Algate, Paul A.

APPLICANT: Fling, Steven P.

TITLE OF INVENTION: COMPOSITIONS AND METHODS

TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF

FILE REFERENCE: 210121.484C5
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                                                                                                                                                                             ; NAME/KEY: misc_feature
; LOCATION: (1)...(1897)
; OTHER INFORMATION: n =
US-09-825-294-214
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 214
LENGTH: 1897
                                                                                                                             Matches 624;
                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: HOMO S
FEATURE:
                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/825,294 CURRENT FILING DATE: 2001-04-03
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CAGCGACAGGCGGCAGCACCAGCCACGAACACCCGGCCGAAACTGCTGCGAGGACAC 180
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                                                                           AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
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                         GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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                                                                                                                                        Similarity
                                                                                                                             Conservative
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                                                                                                                           Score 624; DB Pred. No. 5.9e 0; Mismatches
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5.9e-159;
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                                                                                                                                                  Length 1897;
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APPLICANT: X. Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CF.
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 205
LENGTH: 1619
TYPE: DNA
ORGANISM: Homo sapiens
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Best Local 9
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                                                                                                                                                                                      Local Similarity
                                                                                                                            1 AGTTCTCCTTGCAGAGGGCTGGCGCCGGGACGCGAAGAGCCAACGGGCGCCTGCACAAAGCC 60
CAGCGACAGGCGCAGCACAGCACC-TGCACGAACACCCGGCGAAACTGCTGCGAGGACA 179
                                                                       GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
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                                                      GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 1111
                                                                                                             AGTTCTCCTTGCAGAGGACTGGCGCGGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 1051
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                                                                                                                                                                      624;
                                                                                                                                                                       Conservative
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99.7%;
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                                                                                                                                                                                Score 613; DB 10
Pred. No. 5e-156;
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                                                                                                                                                                                               Length 1619;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 2101-04-03
SOFTWARE: FASTSEQ FOR WINDOWS VERSION SEQ ID NO 211
LENGTH: 1619
TYPE: DNA
ORGANYOW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 211, Application Patent No. US20020004491A1
                                                                                                                                                                                             Matches 624;
                                                                                                                                                                                                           Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHOD
TITLE OF INVENTION: THERAPY AND DIAGNOSIS
FILE REFERENCE: 210121.484C5
                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
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                                                                                                                                                                                                           Local Similarity
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                        CAGCGACAGGCGGCAGCACACCACC-TGCACGAACACCCGCCGAAACTGCTGCGAGGACA 179
                                                                                CCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC 359
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                                                                                                                                      AGTTCTCCTTGCAGAGGACTGGCGCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 1051
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                                                                                                                                                                                              Conservative
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99.7%;
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                                                                                                                                                                                                        Score 613; DB 10
Pred. No. 5e-156;
                                                                                                                                                                                                                                                                                                                                                                                                        AND METHODS FOR THE
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US-09-867-701-2375

Sequence 2375, Application Patent No. US20020132237A1 GENERAL INFORMATION:

US/09867701

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APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REPERENCE: 210121, 497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
           GCCCTCTCGCTCGATGCATGCATGTACCCC--AGCCCTTCTCCTCGTTGCTCTCCTC
                                                                                            GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
                                                                                                                        AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
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Pred. No. 1e-107;
0; Mismatches 14;
                                                                                                                                                                               Indels
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APPLICANT: Jones, Robert

APPLICANT: Harlocker, Susan L.

APPLICANT: Harlocker, Susan L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.497

CURRENT APPLICATION NUMBER: US/09/867,701

CURRENT FILING DATE: 2001-05-29

NUMBER OF SEO ID NOS: 10912

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 4240

TYPE: DNA

TYPE: DNA

TYPE: DNA
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US-09-867-701-4240/c
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Best Local S
Matches 407
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                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapien
170
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                                                                                                           cal Similarity
407; Conserv
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              AAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGT
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 AAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGT
                                                      CATTGATTCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGC
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99.58;
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Pred. No. 1.4e-97;
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\_ \_ Query Match Best Local S Matches 539

Similarity

69.3%;

Conservative

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; OTHER INFORMATION: n - A,T,C US-09-867-701-2375

or,

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NAME/KEY: misc\_feature LOCATION: (1)...(558)

SEQ ID NO 2375 LENGTH: 558

TYPE: DNA

ORGANISM: Homo sapien

FEATURE:

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NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 208
LENGTH: 1362
TYPE: DNA
ORGANISM: Homo sapiens
                                             US-09-867-701-1516/c
Sequence 1516, Application US/09867701
Patent No. US20020132237A1
GENERAL INFORMATION:
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Best Local Similarity 98.9%;
Matches 364; Conservative
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APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN
FILE REFERENCE: 210121.484C5
             APPLICANT: Aglate, Paul A. APPLICANT: Jones, Robert
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CURRENT FILING DATE: 2001-04-03
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Pred. No. 3.5e-88;
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APPLICANT: Jones, Robert
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1532
LENGTH: 390
TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
; TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
; FILE REFERENCE: 210121.497
; CURRENT APPLICATION NUMBER: US/09/867,701
; CURRENT FILING DATE: 2001-05-29
; NUMBER OF SEQ ID NOS: 10912
; SOFTWARE: PastSEQ for Windows Version 4.0
; SEQ ID NO 1516
; SEQ ID NO 1516
; LENGTH: 373
; TYPE: DNA
; ORGANISM: Homo sapien
US-09-867-701-1516
                                              ; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(390)
; OTHER INFORMATION: n =
US-09-867-701-1532
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                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                             Sequence 1532, Application US/09867701 Patent No. US20020132237A1
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches 372;
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Best Local Similarity
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99.7%;
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   Score
Pred.
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Pred. No. 2.6e-88;
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 353; DB 10;
No. 3.8e-86;
                 Length 390;
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517 239 458 179 398 120 338

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Sequence 2409, Application US/09867701

Patent No. US2002013237A1

GENERAL INFORMATION:
APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT APPLICATION SAND CANCER
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2409
LENGTH: 349
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: DNA
; ORGANISM: Homo sapien
US-09-867-701-2409
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US-09-867-701-2409/c
                                                                                                                                                                                                                                                                                                            274 CCAGTCGTGCTTGGGCTTTGGCCGC---AGCCATGATCCTCCGAATCTGGTTGGGCATCCA 330
                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                           GCTTCTTTTGCCAAGCAAGAGAGAGATTTAACACTGTTTCAAACCCGGGGGAGTTGGCT 570
                                                                                CCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAATTTTTCAGTGTTGGACTTGACA 510
                                                                                                                                           AGAAAAGAAAAACACAGGATGAGAAACACAGTAAATGAATAAAACCATAAAATATTTAGCC
                                                                                                                                                                                                                                                                                     CCAGTCGTGCTTGGGGTTTTGCCCCGCAGCCCATGATCCTCCGAATCTGGTTGGGCATCCA 280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAGAAAGACCATTAAATGCTTTAGACAGTGNA 608
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TTTGCCACAAGCAAGAGAGATTTAACACTGTTTCAAACCCGGGGAGTTGGCTGTGTTA
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                                                                                                                          AGAAAAGAAAACACAGCATGAGAACACAGTAAATGAATAAAACCATAAAATATTTAGCC
                                                                                                                                                                                                        GCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAGGAGGGGAGAGACAG
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Pred. No. 8.9e-75;
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APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: COMPOSITIONS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7826
LENGTH: 381
TYPE: DNA
ORGANISM: Homo sapien
US-09-867-701-7826
                                                                       APPLICANT: AGLATE, Paul A.

APPLICANT: Hallocker, Susan L.

APPLICANT: HALLOCKER, SUSAN L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.497

CURRENT APPLICATION NUMBER: US/09/867,701

CURRENT FILING DATE: 2001-05-29

NUMBER OF SEQ ID NOS: 10912

SOFTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 8894

LENGTH: 201

TYPE: DNA

ORGANISM: Homo sapien
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                                                               US-09-867-701-8894
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                                                                                                                                                                                                                                                                                                                                                                 Sequence 8894, Application US/09867701 Patent No. US20020132237A1 GINERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 7826, Application US/09867701 Patent No. US20020132237A1
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Best Local
  Best Local Similarity
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27.4%;
98.3%;
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  Score
Pred.
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  171.2; DB 1
No. 3.3e-37;
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511 159 451 219 391 279 S

331 339 Query Match Best Local Sim Matches 328;

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Sequence 17, Application US/09866582
Patent NO. US20020127620A1
GENERAL INFORMATION:
APPLICANT: Witman, George B.
APPLICANT: Pazour, Gregory J.
APPLICANT: Rosenbaum, Joel L.
APPLICANT: Cole, Douglas G.
TITLE OF INVENTION: INTRAFLAGELLAR TRANSPORT
FILE REFERENCE: 07917-145001
                                                                                                                                                                                 RESULT 15
US-09-866-582-17/c
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; LOCATION: (199)..(1338)
US-09-149-045-1
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US-09-149-045-1/c
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SEQ ID NO 1
LENGTH: 1464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 9.0%;
Best Local Similarity 52.6%;
Matches 123; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: METHODS AND COMPOSITIONS
TITLE OF INVENTION: BETWEEN THE APJ RECEPTOR
FILE REFERENCE: BERLX 73
CURRENT APPLICATION NUMBER: US/09/149,045
CURRENT FILING DATE: 1998-09-08
NUMBER OF SEQ ID NOS: 8
SOFTWARE PROPERTY OF SEC ID NOS: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1, Application US/09149045 Patent No. US20020062488A1 GENERAL INFORMATION:
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APPLICANT: HORUK, RICHARD
APPLICANT: ZHOU, YIQING
APPLICANT: HESSELGESSER, JOE
APPLICANT: DOMS, ROBERT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1182 CTGGCCACAGCAGAGCATGGAGGTGCAGGCCTGGCGGAAGCGGGGGTCGAAAAAGGCATA 1123
                                                                                                                                                                                                                                                         1002 CTTCACCAGGTGGTAGGGCATCCAGCACAGGGCAAAGGTCACCACCAGCACCAC 949
                                                                                                                                                                                                                                                                                                                                   1062 CATGAGGAAGAGGTCAAAGTCACAGGGCCAGTGCAGCAGGCTGCCCAGCATGTACAGCGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  127 CAGGCGGCAGCACCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACACCGTGTA 186
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61
                                                                                                                                                                                                                                                                               CCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC 360
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US-09-866-582-17
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CURRENT FILING DATE: 2001-05-24
PRIOR APPLICATION NUMBER: US 60/206,923
PRIOR FILING DATE: 2000-05-24
NUMBER OF SEQ ID NOS: 50
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 17
LENGTH: 2036
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Best Local (
                                                                                                                                                                                                                                                                                                                                        Best Local Similarity Matches 103; Conserv
time : 34.1619
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Chlamydomonas reinhardtii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
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                                                                                                                                 422 CAGCTCCGTGAAGCGCTCCAGCGCGGAGCGGTCCGAGGGCGCCATGTTGGAGGGCGGGTT
                                                                 362 CTTGATGCTCTTGAG 348
                                                                                                 199 GATGACCGAGCTGAG 213
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1: /cgn2_6/ptodata/1,
2: /cgn2_6/ptodata/1,
3: /cgn2_6/ptodata/1
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625
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/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
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| 39               | 39               | 39               | 39               | 39               | 39               | 39               | 39               | 39               | 39               | 39               | 39               | 39               | 39               | 39              | 39              | 39               | 39               |
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| US-08-459-504B-4 | US-08-459-504B-2 | US-08-459-595A-8 | US-08-459-595A-4 | US-08-459-595A-4 | US-08-459-595A-2 | US-08-459-448A-8 | US-08-459-448A-4 | US-08-459-448A-4 | US-08-459-448A-2 | US-07-951-715A-8 | US-07-951-715A-4 | US-07-951-715A-4 | US-07-951-715A-2 | US-09-547-422-3 | US-08-459-444-3 | US-08-459-504B-3 | US-08-459-595A-3 |
| Sequence 4,      | Sequence 2,      | Sequence 8,      | Sequence 4,      | Sequence 4,      | Sequence 2,      | Sequence 8,      | Sequence 4,      | Sequence 4,      | •                | Sequence 8,      | Sequence 4,      | Sequence 4,      | Sequence 2,      | Sequence 3,     | Sequence 3,     | Sequence 3,      | Sequence 3,      |
| Appli            | Appli            | Appli            | Appli            | Appli            | Appli            | Appli            | Appli            | Appli            | Appli            | Appli            | Appli            | Appli            | Appli            | Appli           | Appli           | Appli            | Appli            |
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## ALIGNMENTS

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RESULT 1
US-08-153-848-39/c
US-08-153-848-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 5759804 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                       APPLICATION UMBER: US/08/153
EILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US 07/977
FILING DATE: 17.NOV-1992
ATTORNEY/AGENT INFORMATION:
NAME: NO. 5759804and, Greta E
NAME: NO. 5759804and, Greta E
REGISTRATION UMBER: 35,302
REFERENCE/DOCKET NUMBER: 3179
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                               TELEPHONE: (312) 474-6300
TELEFAX: (312) 474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 39
                FEATURE:
NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Godiska, Ronald
APPLICANT: Gray, Patrick W.
APPLICANT: Schweikart, Vicki L.
TITLE OF INVENTION: NO. 5759804el Seven Transmembrane Receptors
NUMBER OF SEQUENCES: 64
CORRESPONDENCE ADDRESS:
                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 1872 base pairs
                                                                           MOLECULE TYPE:
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ZIP: 60606
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CITY: Chicago
STATE: Illinois
                                                                                                               STRANDEDNESS:
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                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
              MOLECULE TYPE:
                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                             TELEPHONE: (312) 474-6300
                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/977,452
FILING DATE: 17-NOV-1992
FILING DATE: 11-NOV-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 09/088,337 FILING DATE: 01-JUN-1998 PRIOR APPLICATION DATA:
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FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Godiska, Ronald
APPLICANT: Gray, Patrick W.
APPLICANT: Schweikart, Vicki L.
TITLE OF INVENTION: No. 6107475el Seven Transmembrane Receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
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                                 TOPOLOGY:
                                                                                                                                                                                                            NAME: Jill E. Uhl
REGISTRATION NUMBER: 43,213
REFERENCE/DOCKET NUMBER: 27
                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CITY: Chicago
                                                                                                                                                            TELEFAX:
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                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 0: FILING DATE: 17-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PatentIn Release #1.0, Version #1.25
                                   linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Borun
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             DNA (genomic)
                                                single
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52.6%;
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Best Local Similarity
Matches 123; Conserv
                                                                                                                                        TELEFAX: (312) 474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 39:
                                                                                                                                                                         CLASSIFICATION.

PRIOR APPLICATION UMBER: US 08/153,848

FILING DATE: 17-NOV-1993

APPLICATION NUMBER: US 07/977,452

FILING DATE: 17-NOV-1992

ATTORNEY/AGENT INFORMATION:

NAME: NO. 6348574and, Greta E.

REGISTRATION NUMBER: 35,302

REGISTRATION NUMBER: 35,302

REGISTRATION NUMBER: 31794

TELECOMMUNICATION INFORMATION:

TELEPHONE: (312) 474-6300
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LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/088,337B
FILING DATE: 01-Jun-1998
                 MOLECULE
                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: NO. NUMBER OF SEQUENCES: 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Godiska, Ronald
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTTCACCAGGTGGTAGGGCATCCAGCACAGGGCAAAGGTCACCACCAGCACCAC 952
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAAAATCAGCCC 360
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                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                  LENGTH: 1872 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 6300 Sears Tower, CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Marshall, O'Toole, Gerstein, Murray &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
              TYPE: DNA (genomic)
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Schweikart, Vicki L.
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                                    linear
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Pred. No. 5.3e-06;
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; NAME/KEY: CDS; LOCATION: 202..1341; SEQUENCE DESCRIPTION: SEQ ID NO: US-09-088-337B-39
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                                                                                                  TELEFAX: (312) 474-044
TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1872 base pair
                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/977,452
FILING DATE: 17-NOV-1992
ATTORNEY/AGENT INFORMATION:
NAME: NOLAND, Greta E.
REGISTRATION NUMBER: 35,302
REFERENCE/DOCKET NUMBER: 31794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
TELEFAX: (312) 474-0448
                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                  FEATURE:
                               MOLECULE TYPE: DNA (genomic)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
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                                                               LENGTH: 1872 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE:
NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
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                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Marshall, O'Toole, Gerstein,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGAT 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      60606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schweikart, Vicki L.
WENTION: Novel Seven Transmembrane Receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gray, Patrick W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Godiska, Ronald
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                                                   linear
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52.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                             Version #1.25
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9:
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US-08-499-215-1/c
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Patent No. 5612204
GENERAL INFORMATION:
APPLICANT: Saeki, Hisashi
APPLICANT: Miura, Akira
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 123;
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                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 6379 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, V SOFTWARE: & WordPerfect version 5. CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPUTED:
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                           HYPOTHETICAL: NO ORIGINAL SOURCE:
                                                                                                                                                                                                            TOPOLOGE TYPE: DN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1185 CTGGCCACAGCAGAGCATGGAGGTGCAGGCCTGGCGGAAAGCGGGGGGGTCGAAAAAAGGCATA 1126
                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1005 CTTCACCAGGTGGTAGGGCATCCAGCACAGGGCAAAGGTCACCACCAGCACCAC 952
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FEATURE:
                                                                      FEATURE:
                                                                                                                                             FEATURE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: BIOLOGICAL DEGRADATIVE
TITLE OF INVENTION: TREATMENT OF CHLORINE-SUBSTITUTED ETHYLENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        307 CCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAATCAGCCC 360
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     127 CAGGCGGCAGCACCACCACGAACACCCGGCCGAAACTGCTGCGAGGACACCGTGTA 186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            247 CATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTCGTTGGGTTTGGCCGCAGCCATGAT 306
                                 NAME/KEY:
LOCATION:
                                                                                   NAME/KEY: CDS
LOCATION: 910..1935
OTHER INFORMATION:
                                                                                                                                                            STRAIN:
                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: JP Hei-6-179689 FILING DATE: 08-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 21
CITY: N.W.
                  OTHER INFORMATION:
                                                                                                                                                                              STRAIN:
                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
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                                                                                                                                                                                                                                                            nucleic acid
DEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Washington D.C
                                                                                                                                                    No. 5612204ardia corallina B-276 (FERM P-4094; FERM BP-5124; ATCC 31338)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2: Sughrue, Mion,
2100 Pennsylvania
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                                                     CDS
                                                                                                                                                                                                                                                 linear
                                 1935..2285
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                /product= "amoB"
                                                                                     /product=
                                                                                                                                                                                                                                                                                                                                                                                                                                                 US/08/499,215
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Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 56.4; DB 5; Pred. No. 5.3e-06;
                                                                                       "amoA'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches 111;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Version #1.30
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OTHER INFORMATION: /pr
; FEATURE;
; NAME/KEY: CDS
; LOCATION: 3805..4830
; OTHER INFORMATION: /pr
US-08-499-215-1

/product= "amoD"

LOCATION: NAME/KEY:

2300..3802

/product= "amoC"

Query Match

Local Similarity

8.0%;

Score 50; DB 1; Pred. No. 0.00054;

Matches

125;

Conservative

0;

Mismatches

125;

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118

178

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AGCCATGATC 307

1293

238

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1233 CCCCATGGTC 1224

US-08-232-463-14/c

Sequence 14, Application US/08232463 Patent No. 5670367

APPLICANT: DORNER, F.
APPLICANT: SCHEIFLINGER, F.
APPLICANT: FALKMER, F. G.
TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS
NUMBER OF SEQUENCES: 52

CORRESPONDENCE ADDRESS:

GENERAL

INFORMATION:

APPLICATION NUMBER: EP 91 114
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 3047
TELECOMMUNICATION INFORMATION:

JMBER: EP 91 114 300.6 26-AUG-1991

US/07/935,313

TELEFAX: TELEPHONE:

(703)683-4109

(703)836-9300

30472/114 IMMU

APPLICATION NUMBER: US FILING DATE: CLASSIFICATION: 435 PRIOR APPLICATION DATA: APPLICATION NUMBER: US

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:

Version

US/08/232,463

COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk

22313-0299 ٧A

COUNTRY:

USA

CITY: Alexandria

STREET:

ADDRESSEE:

E: Foley & Lardner 1800 Diagonal Road,

Suite 500

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1413 GTCCTGGAGCAGCCGCATGCGGTCCACGGCCTGGAACACCACGCTGAACTGGACCGTGTC 1354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1473 GCGGGCCCCGGCGTGAATTCCGGCGACTCCTGCAGGTGGTGCTAGGTGGTGGACGAT 1414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              58 GCGGGCGCTGTGGTGGAGTGCGCATGTACGCGCAGGCGCCTTCTCGTGGTTGGCGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCAGGCGGCGTAGGACCTGGCGAGGATCTTCTGCGACCAGATGGGCGTGATCGCCGCCGA 1234
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGG 237
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GGACATGGCCTGGCGCACGGCGTAGGCCCAGGCTCAGGAAGAGCCCCGTACTCGACGAATGG
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                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/200,090
CURRENT FILING DATE: 1998-11-25
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 3
LENGTH: 1176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence 3, Application US/09200090
Patent No. 6420137
                                                                                                                                   Matches 124;
                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches 11;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Strand, Joann
APPLICANT: Hadcock, John R.
APPLICANT: Hadcock, John R.
TITLE OF INVENTION: No. 6420137el cDNA Encoding The Long and
TITLE OF INVENTION: Forms of The Human Neurotensin Subtype 2
FILE REFERENCE: 0630/0E792
                                                                                                                                                                                                                               TYPE: DNA ORGANISM: human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1134
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                                                          1090 TGGGGTGGTGCTCTCCACACAGGGAGCTGACGGCTTCCAGGAAGAGTTTTCTGAAGGAGG 1031
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IMMEDIATE SOURCE:
CLONE: pTZgpt-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            193
171 GCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCG
                                                                                            111 TGGCGTGCTGCAGCGACAGGCGGCAGCACCAGCACCTGCACGAACCCCGCCGAAACTGCT 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             373
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                                                                                                                                   Conservative
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                                                                                                                                   0;
                                                                                                                                                     Score 46.6; DB Pred. No. 0.002;
                                                                                                                                   Mismatches 129;
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                                                                                                                                                                     DB 4;
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                                                                                                                                                                      Length 1176;
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CURRENT APPLICATION NUMBER: US/09/007,005B
CURRENT FILING DATE: 1998-01-14
EARLIER APPLICATION NUMBER: 60/035,963
EARLIER APPLICATION NUMBER: 60/064,491
EARLIER APPLICATION NUMBER: 60/064,491
EARLIER FILING DATE: 1997-11-06
NUMBER OF SEQ ID NOS: 33
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 17
LENGTH: 289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; NAME/KEY: misc_feature
; LOCATION: (1)...(289)
; OTHER INFORMATION: n = A,T,C or G
US-09-007-005-17
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RESULT 9
US-09-244-796-17
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APPLICANT: SZOSTAK, Jack W.
APPLICANT: ROBERTS, RICHARD W.
APPLICANT: Liu, Rihe
TITLE OF INVENTION: SELECTION OF PROTEINS USING RNA-PROTEIN
TITLE OF INVENTION: FUSIONS
FILE REFERENCE: 00786/350003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17, Appl. Patent No. 6258558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 28
TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Translation template
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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                                                                                                                                                                                                                                                                                                              96
                                                                                                                                                                                                                                                                                                                                                                                       36 CRARARURURARCRARARURGRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNR 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                 AACACA 406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CGATGGCTCCAGG 838
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                                                                                                                                                                                          AATGTCACAACAATCAGCCCTGGGCAGACACGAGCAGGAGGGGGGAGAGACAGAGAAAAGAAA 400
                                                                                                                                                                                                                                 NRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNR
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                                                                             AAAAAA 281
                                                                                                                                                      TGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCC
                                                                                                                                                                                                                                                                                                            NRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ch 7.4%; Score 46; DB 4; Length 289, 1 Similarity 9.3%; Pred. No. 0.0015; 23; Conservative 103; Mismatches 120; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/09007005B
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; LOCATION: (1)...(289)
; OTHER INFORMATION: n = A,T,C
US-09-244-796-17
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CURRENT FILING DATE: 1999-02-05
EARLIER APPLICATION NUMBER: 60/035,963
EARLIER FILING DATE: 1997-01-27
EARLIER APPLICATION NUMBER: 60/064,491
EARLIER FILING DATE: 1997-11-06
EARLIER FILING DATE: 1997-11-06
EARLIER APPLICATION NUMBER: 09/007,005
EARLIER FILING DATE: 1998-01-14
NUMBER OF SEQ ID NOS: 33
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                         US-08-832-399-1/c
                                                                                                                                                                                                                                                                                                                                               RESULT 10
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                                                                                                                                                                                                                                                        Sequence 1, Application US/08832399 Patent No. 6008050 GENERAL INFORMATION:
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Best Local Similarity
Matches 23; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: SELECTION OF PROTEINS USING TITLE OF INVENTION: FUSIONS FILE REFERENCE: 00786/350007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Szostak, Jack W. APPLICANT: Roberts, Richard W. APPLICANT: Liu, Rihe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 289
TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                        NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation
COMPUTER READABLE FORM
                                                                                                                                                                                         APPLICANT: Shabon, Usman TITLE OF INVENTION: NOVE
                                                                                                                                                                                                                                      APPLICANT:
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                                                              STREET: 709 Swedeland CITY: King of Prussia STATE: PA
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                        ZIP:
                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         36 CRARARURURARCRARARURGRNRNRSRNRNRSRNRNRSRNRNRSRNRNRSRNR 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCG 280
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                      19406
                                                                                                                                                                                                                                    Bergsma,
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9.3%; Pred. No. 0.0015;
                                                                                                                                                                                         NOVEL HUMAN NEUROTENSIN RECEPTOR TYPE
                                                                                                         Road
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GENERAL INFORMATION:
APPLICANT: Derk J. Bergsma
APPLICANT: Usman Shabon
TITLE OF INVENTION: NOVEL HUMAN NEUROTENSIN RECEPTOR TYPE
TITLE OF INVENTION: AND SPLICE VARIANTS THEROF
FILE REFERENCE: GH-50020-1
CURRENT APPLICATION NUMBER: US/09/372,498
CURRENT FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: 08/832,399
PRIOR FILING DATE: 1997-04-02
NUMBER OF SEQ ID NOS: 12
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SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 45.6; DB
Pred. No. 0.004;
0; Mismatches 1
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Best Local Similarity 49.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/08858876A Patent No. 6022856
                                                                                                                                                                                                                           NAME: Player, William E.
REGISTRATION NUMBER: 31,049
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                   CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/FF
FILING DATE: 17-MAR-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL
                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/858,876A
FILING DATE: 19-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                        FEATURE:
                                                                               MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 1575 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Type 2 Neurotensin Receptor TITLE OF INVENTION: (hNT-R2) NUMBER OF SEQUENCES: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Daniel CAPUT
APPLICANT: Pascale CHALON
APPLICANT: Pascual FERRARA
APPLICANT: Vita NATALIO
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NAME/KEY:
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                                                            ORGANISM:
                                                                                                                          TOPOLOGY:
                                                                                                                                         STRANDEDNESS: single
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                                                            Homo sapiens
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; MOLECULE TYPE: CDNA US-08-832-399-1

TOPOLOGY:

linear single INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 1342 base pairs

TYPE: nucleic acid STRANDEDNESS: sing

ATTORNEY/AGENT INFORMATION:
NAME: King, William T
REGISTRATION NUMBER: 30,954
REFERENCE/DOCKET NUMBER: GHSG
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-770-5515

GH50020

610-270-5090

PRIOR APPLICATION DATA:

FILING DATE: APPLICATION NUMBER: APPLICATION NUMBER: US/08/832,399 FILING DATE: 02-APR-1997 CLASSIFICATION: 435

OPERATING SYSTEM: SOFTWARE: FastSE(

TYPE:

IBM Compatible

Best Loc Matches

Local Similarity les 120; Conserv

Conservative

7.3%;

171

Query Match

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351 CAAT 354

746

686 CGAT 683

RESULT 11 US-09-372-498-1/c

Sequence 1, Application US/09372498 Patent No. 6166182

SOFTWARE: SEQ ID NO 1

FastSEQ for Windows Version 3.0

LENGTH: 1342

TYPE: DNA
ORGANISM: HOMO SAPIENS

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US-09-472-880-1/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches 120;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 7.3%;
Best Local Similarity 49.2%;
                                                                                                                                                                REGISTRATION NUMBER: 31,049
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 1575 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1123
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        943 CGAT 940
                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/472,880
FILING DATE: 28-Dec-1999
CLASSIFICATION: <UNknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 20004

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Daniel CAPUT
Pascale CHALON
Pascual FERRARA
                                                                                                                                                                                                                                              APPLICATION NUMBER: PCT/FR 9723204
FILING DATE: 17-MAR-1997
ATTORNEY/AGENT INFORMATION:
NAME: Player, William E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Jacobson, Price, Holman & Stern,
STREET: 400 Seventh Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Type 2 Neurotensin Receptor (hNT-R2)
NUMBER OF SEQUENCES: 12
                                         FEATURE:
                                                                           ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CAAT 354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TCACCATGTAGAAGTAGTAGAAAATTGTACAGTGGGTCAGTCCACGCGTCATCAGGTA 1004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGGACACGGCGTTGTAGAGAAGAGGAGTCACAGCTGAGCTGACGTAGAAAAGTGTGTTGG 1064
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                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Washington STATE: D.C.
                                                                                           TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
                   NAME/KEY:
                                                         ORGANISM: Homo sapiens
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CDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Strand, Joann
APPLICANT: Hadcock, John R.
APPLICANT: Hadcock, John R.
TITLE OF INVENTION: No. 6420137el cDNA Encoding The Long and Short
TITLE OF INVENTION: Forms of The Human Neurotensin Subtype 2 Receptor
FILE REFERENCE: 0630/0E792
CURRENT APPLICATION NUMBER: US/09/200,090
CURRENT FILING DATE: 1998-11-25
NUMBER OF SEQ ID NOS: 18
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1
LENGTH: 1233
TYDE: DNA
ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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Best Local Similarity
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Best Local Similarity 48.8%;
Matches 119; Conservative
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907 CGAT 904
                                        351
                                                                                 967
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                                        CAAT 354
                                                                             AGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTT 290
                                                                                                                                                                                                                                                                                                                             TGGGGTGGTGCTCCACACAGGGAGCTGACGGCTTCCAGGAAGAGTTTTCTGAAGGAGG 1088
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                                                                                                                 TGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAA 350
                                                                                                                                                          TCACCATGTAGAAGTAGTGGTAGAAATTGTACAGTGGGTCAGTCCACGCGTCATCAGGTA
                                                                                                                                                                                                                                            AGGACACGGCGTTGTAGAGAAGAGGAGTCACAGCTGAGCTGACGTAGAAAAGTATGTTGG 1028
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                                                                                                                                                                                                                                                                                                                                                                                                          Score 44; DB 4; Length 1233
Pred. No. 0.011;
0; Mismatches 125; Indels
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Pred. No. 0.0044;
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RESULT 15 US-09-668-680-12/c

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NUMBER OF SEQ ID NOS: 13
SOFTWARE: pt_FL_genes Version 2.0
SEQ ID NO 12
LENGTH: 1535
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: (1)..(1338)
US-09-668-680-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Tang, Y. Tom
APPLICANT: Liu, Chenghua
APPLICANT: Zhou, Ping
APPLICANT: Asundi, Vinod
APPLICANT: Asundi, Vinod
APPLICANT: Wang, Jian-Rui
APPLICANT: Wang, Jian-Rui
APPLICANT: Wang, Jian-Rui
APPLICANT: Xue, Aidong J.
APPLICANT: Xue, Chongjun
APPLICANT: Drmanac, Radoje T.
APPLICANT: Drmanac, No. 6436703el Nucleic Acids and
TITLE OF INVENTION: Polypeptides
FILE REFERENCE: 790CIP2A
CURRENT APPLICATION NUMBER: US/09/668,680
CURRENT FILING DATE: 2000-09-22
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR APPLICATION NUMBER: 09/640,217
PRIOR APPLICATION NUMBER: 09/640,217
PRIOR FILING DATE: 2000-03-31
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Search completed: November Job time: 39.4202 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 12, Application US/09668680 Patent No. 6436703 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               1052 AGG 1050
                                                                                                                                                                                                                                                                            1172 CACAGGGCCTCCTGGAAGGTCTCTCGGAAGCGGCTGGACATGAGGCTATAGAGCACGGGG 1113
                                                                                                                                                                                                                                                                                                                                                                       1112 TTGGCCGCCGAGCCCAGGTAGAAGAAGAAGATGCCGGAGATGACGTGCACGTGCTGGAAGGCC 1053
                                                                                                                                                                                                                                                                                                       137 CACAGCACCTGCACGAACACCCGCCGAAAACTGCTGCGAGGACACCCGTGTACAGGAGGGGG 196
                                                                                                                                                                                                          197 TTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGAGGATCATGTACGCC
                                                                                                                                      257 CGG 259
                                                                                                                                                                                                                                                                                                                                                                                                                77 GAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGGCAG 136
                   7, 2002, 15:58:19
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Maximum DB seq length: 2000000000
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1: gb_ba:*
2: gb_htg:
3: gb_in:*
4: gb_om:*
 US-09-970-966-210
625
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
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Copyright (c) 1993 - 2002 Compugen Ltd.
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gb_ph:*
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Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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|------------|-------|----------------|----------------|----------|-------------|---------------|-----------|--------|------|------|-----|-----|-----|--------------------|------------|--------------|-----------|------------|--------------|--------------|---------|-----|-----|--------------------|------------|---------------|--------|---------------|----------|--------------|-------------|-----------------|------------|--------------|---------|--------------|---------|---------|---------------|---------------|-------------|---------------|---------------|--|
|            | 4     | 40.2           |                |          | # /         | 47            | 47.       | 47.    | 48.  | 49.  | ( n | (,, | (n  | cn e               | ,<br>,     | л U<br>- С   | n 0       | , ,        | 56.          | 56.          | 56.     | 56. | 56. | 56.4               | л (<br>Л ( | л U           | , v    | 59.           | _14      | 230.         | 300.        | 30              | 361.       | 361.         | 3 3 9   | 39           | 412.    | 479.    | 598.          | on.           |             | 6             | 8             |  |
|            | 7.4   | 4              | ٠.             | ٠        | 10          | ·<br>0        |           |        | . 7  | 9    | ٠   |     |     |                    |            | o c<br>⊢     |           |            | •            |              |         |     |     | ۰.<br>٥٠           |            |               | •      | •             | Ñ        | 6.9          | 8.1         | 8 :<br>3 :<br>2 | •          | ٠,           | Ņ       | Ñ            | ō       | 6.      | 95.7          | 6.7           | .7          | 9             | ch L          |  |
|            | ٧     | 000            | 000            | 0000     | 100         | 7929          | 9437      | 5362   | 8050 | 7218 | 379 | 379 | 379 | 379                | 270        | 20/4<br>44/4 | 9216      | 1872       | 872          | 801          | 754     | 583 | 148 | 1140               | 140        | 121           | 104    | 917           | 0503     | 1797         | 2874        | 9885            | 508        | 362          | 524     | 524          | **      | 587     | 890           | 1832          | 9676        | 826           | ngth D        |  |
| ALIGNMENTS | AKIDA | AF 4 2 9       | ACLO           | SC3A/    | BCOZZOO     | 2 AC128436    | AC10341   | SCF56  |      |      |     |     |     | 6 E08269           | <b>v</b> : | > 2          | A         | Α          | >            | Ξ            | B       | Ξ   | ا ح | 6 E63120           | 7 0        | 9             | E11480 | AB0170        | AC128363 | AB0416       | AC11207     | AC12449         | 6 AF034033 | AX31994      | AX36225 | AX35876      | AX15012 | AX13669 | AX13628       | AK09450       | AC07977     | BC01144       | Ü             |  |
| S          | 69070 | 429313 HOHO Sa | 100744 OFYZA S | Streptom | OZZZZZ HOMO | 128436 Rattus | 3419 Ratt |        |      |      |     |     |     | E08269 qDNA of alk |            |              | 1786 Homo | 1356 Seque | 7256 Sequenc | 71 H.sapiens | 2688 нс | 2   | 2   | E63120 Peptide der | 50         | 70 u sanience | 5      | 17027 Mus mus | 3363 R   | 11649 Mus mu | 2072 Rattus | 493             | 1000       | 942 Sequence | 62255 S | 3762 Sequenc | 50120 S | 36698 S | 36281 Sequenc | 1501 Ното зар | 079773 Homo | 1449 Ното зар | escription    |  |

| JOURNAL                                                          | TITLE             | AUTHORS        | REFERENCE          |                                                            |                                                                   | ORGANISM     | SOURCE        | KEYWORDS | VERSION                | ACCESSION | DEFINITION                               | LOCUS                                        | BC011449 | RESULT 1 |
|------------------------------------------------------------------|-------------------|----------------|--------------------|------------------------------------------------------------|-------------------------------------------------------------------|--------------|---------------|----------|------------------------|-----------|------------------------------------------|----------------------------------------------|----------|----------|
| Submitted (25-JUL-2001) National Institutes of Health, Mammalian | Direct Submission | Strausberg, R. | 1 (bases 1 to 826) | Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. | Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; | Homo sapiens | Homo sapiens. |          | BC011449.1 GI:15277472 | BC011449  | Homo sapiens, clone IMAGE:4026092, mRNA. | BC0111449 826 bp mRNA linear PRI 22-AUG-2001 |          |          |

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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Institute for Systems Biology
http://www.systemsbiology.org
contact: amadan@systemsbiology.org
Anup Madan, Rachel Dickhoff, Jessica Fahey, Stephanie For
Greene, Mark Ketteman and Anuradha Madan
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Tissue Procurement: ATCC/DCTD/DTP
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Contact: MGC help desk
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/clone_lib="NIH_MGC_20"
/lab_host="DH10B-R"
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/db_xref="taxon:9606"
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197 c 228 g
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Submitted (09-AUG-2001) Genome Sequencing Center, Washington
Statement School of Medicine, 4444 Forest Park Parkway, St.
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                                                                                                   Submitted (09-JAN-2002) Department of Genetics, Washington University, 4444 Forest Park Avenue, St. Louis, Missouri 63108, USA On Aug 9, 2001 this sequence version replaced g1:14488388.
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Shah,N., Meyer,R., Boyer,E. a
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 129676)
Sulston J.E. and Waterston,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2001)
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            Web site: http://genome.wustl.edu/gsc
Contact: sapiens@watson.wustl.edu
------Summary Statistics
Center project name: H_NH0258B17
                                                                Center: Washington University Genome Sequencing Center Center code: WUGSC
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NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap between neighboring data submissions.

all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >-30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by restriction This sequence was finished as follows unless otherwise noted: digest.

Mapping Mapping INFORMATION:
information for this clone was provided
on, Department of Genetics, Washington Ur University, γd Dr. St. Ö. Louis

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NEIGHBORING SEQUENCE INFORMATION:
The clone sequenced to the left is RP11-1L22; the clone sequenced to the right is RP11-159N20, 2000 bp overlap. Actual start of this clone is at base position 1 of RP11-258B17; actual end is at base position 60003 of RP11-159N20.
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/rpt_family="(TG)n"
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/db_xref="taxon:9606"
/chromosome="2"
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35361. .35676
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                                                    Score 607; DB 9; Pred. No. 7.5e-129; O; Mismatches 1;
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FEATURES

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Direct Submission

Submitted (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team); 2-6-7

Kazusa-Kamatari, Kisarazu, Chiba 292-0812, Japan

(E-mail:genomics@hri.co.jp, Tel:81-438-52-3975, Fax:81-438-52-3986)

NEDO human cDNA sequencing project supported by Ministry of
                                                                                                                                                   Kawakami, B., Sugiyama, A., Janessovo, T., Yamamoto, J., Isono Otsuki, T., Sato, H., Wakamatsu, A., Ishii, S., Yamamoto, J., Isono Kawai-Hio, Y., Saito, K., Nishikawa, T., Kimura, K., Yamashita, H., Kanda, K., Nakamura, Y., Sekine, M., Kikuchi, H., Kanda, K., Matsuo, K., Nakamura, Y., Sekine, M., Kikuchi, H., Kanda, K., Wagaisuma, M., Murakawa, K., Kanehori, K., Takahashi-Fujii, A., Wagaisuma, M., Murakawa, K., Kanehori, K., Magaisuma, M., Murakawa, K., Kanehori, K., Magaisuma, M., Murakawa, K., Kanehori, K., Masuho, Y., Nagai
                                                                                                                                                                                                                                                                                           Homo
                                                                                                                                                                                                                                                                                                                     AK094501.1 GI:21753575 oligo capping; fis (full insert sequence). Homo sapiens alzheimer cortex cDNA to mRNA,
                                                                                                          Unpublished
                                                                                                                                          and Isogai,T.
                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae
                                                                                                                                                                                                                                                                                                          clone:BRALZ2001350.
                                                                                                                                                                                                                                                                                                                                                                                                to Homo
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TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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605; Conservative
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NCTVNVQDMCQKEVMEQSAGIMYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPLCNGPRPKKRGSSASALRPGLRTTILFLKLALFSAHC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone="BRALZ2001350"
/tissue_type="alzheimer
/clone_lib="BRALZ2"
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/db_xref="GI:21753576"
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/db_xref="taxon:9606"
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AX136281
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AATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACC
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Patent: EP 1067182-A 203 10-JAN-2001;
Helix Research Institute (JP)
Location/Qualifiers
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/db_xref="GI:14272688"
/db_xref="GI:14272688"
/translation="mwtlGIAATFCGLFLLPGFALQIQCYQCEEFQLNNDCSSPEFIV
NCTVNVQDMCQKEVMEQSAGINYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPLCNQPRPKKRGSSASALRPGLRTTILFLKLASSRHTAELKEMPPPPALFFQPSPPT
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/db_xref="taxon:9606"
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                                               AGCTTCTTTTGCCACAAGCAAGAGAGAATTTAACACTGTTTCAAACCCGGGGGAGTTGGC
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AX136698
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Patent: EP 1067182-A 620 10-JAN-2001;
Helix Research Institute (JP)
Location/Qualifiers
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Pred. No. 8.8e-100;
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Kroes,R.A., Moskal,J.R. and Yamamoto,H.
Differential gene expression in cancer
Patent: WO 0136885-A 95 25-MAY-2001;
NYXIS NeuroTherapies, Inc. (US)
Location/Qualifiers
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AX150120
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Sequence 15 from
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                                 Baker, K.P., Desnoyers, L., Gerritsen, M.E., Goddard, A., Godowski, P.J., Grimaldi, J.C., Gurney, A.L., Smith, V., Watanabe, C.K. and Wood, W.I.
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                       Secreted and transmembrane polypeptides
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/db_xref="taxon:9606"
113 c 94 g 14:
 0193983-A 15
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WO0193983
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         Query Match
Best Local Similarity
Matches
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                                                                                                               Genentech,
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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15 from Patent WO0208288
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/db_xref="taxon:9606"
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Location/Qualifiers
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/db_xref="taxon:9606"
433 c 435 g 33
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Eukaryota;
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Sequence 1 from Patent WO0181634.
AX319942
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Methods and compositions for the diagnosis and treatment cardiovascular and tumorigenic disease using 4941
Patent: WO 0181634-A 1 01-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                             Millennium Pharmaceuticals, Inc. (US)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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LLGNSATIRVTQVLQKKGYLQKEVTDHHVSLACSDILVFLIGMPWEFYSIIWNPLTTS
SYTLSCKLHTELFERGYSATILHVLTILSFEWI JAICHPERYKAVSGFQUYKLIGFWW
VTSALVALPLLFAMGTEYPLVKVPSHRGITCNRSSTRHHEQPETSNNSICTNLSSRWT
VFQSSIFGAFVVYLVSVAFMCWNMGVLMKSQKGSLAGFTRPPQLRKSBSEESRT
ARROTIIFLKIVVLLAVCMMPNQIRRIMAANFKHDWIRSYFRAYMILLPESETFFY
LSSVINPLLYTVSSQOFRRVFYQVLCCRLSLQHANHEKRLEVHAHSTIDSAREVQRPL
LEASRQSSARRTEKIFLSTFQSEAEPQSKSQSLSLESLEPNSGAKPANSAAENGFQE
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/protein_id="CAD19381.1"
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/db_xref="taxon:9606"
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1 (bases 1 to 1362)

McKee, K. K., Tan, C.P., Palyha, O.C., Liu, J., Feighner, S.D., Hreniuk, D.L., Smith, R.G., Howard, A.D. and Van der Ploeg, L.H. Cloning and characterization of two human G protein-coupled receptor genes (GPR38 and GPR39) related to the growth hormone secretagogue and neurotensin receptors

Genomics 46 (3), 426-434 (1997)
                                                                                                                                                                                                                                                                                                    Direct Submission
Submitted (17-NOV-1997) Biochemistry and
Inc., PO Box 2000, Rahway, NJ 07065, USA
Location/Qualifiers
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McKee, K.K., Tan, C.P., Palyha, O.C.,
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Sequence 3
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
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Pred. No. 1.1e-72;
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FEATURES

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SOURCE KEYWORDS VERSION ACCESSION

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TGGGCAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC
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                                                                                                                                                                                                                                                                                                                                                   AC124493
AC124493.2 GI:2169
HTG; HTGS_PHASE1; H
house mouse.
                                                                                                                           Direct Submission
Submitted (14-JUN-2002) Genome Sec
Parkway, St. Louis, MO 63108, USA
3 (bases 1 to 209885)
McPherson, J. D. and Waterston, R. H.
Direct Submission
Center: Washington University Genome Sequencing Center code: WUGSC
Web site:http://genome.wustl.edu/gsc/index.shtml
Contact: submissions@watson.wustl.edu
                                                                                    Submitted (05-JUL-2002) Genome Sequencing Parkway, St. Louis, MO 63108, USA On Jul 5, 2002 this sequence version repla
                                                                                                                                                                                                                                                                                                                                                                                                     AC124493 209885 bp DNA LINE Mus musculus chromosome UNK clone RP23-462P13, SEQUENCE, 5 unordered pieces.
                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                     Mus musculus
                                                                                                                                                                                                                      McPherson,J.D.
                                                                                                                                                                                                                                                Unpublished
                                                                                                                                                                                                                                                               The sequence of Mus musculus
                                                                                                                                                                                                                                                                               McPherson, J.D. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity 98.964; Conservative
                                                                                                                                                                                                                                 (bases 1
                                                                                                                                                                                                                                                                                           (bases 1 to 209885)
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                                                                                                                                                                                                                                                                                                                                                            GI:21699722
HASE1; HTGS_DRAFT; HTGS_FULLTOP
                                                                                                                                                                                                                                  to 209885)
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98
                                                                                                                                                                                                                     and Waterston, R.H.
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                                                          Genome Center ----
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s musculus clone
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Pred.
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Sciurognathi; Muridae;
                                                                                                                                                                                        Sequencing
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                                                                                                                                                                                          4444
                                                                                                                   4444 Forest
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                                                                                                                                                                                                                                                                                                          Murinae;
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                                                                                                                                                                                          Forest
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                                                                                                                                                                                          Park
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BASE COUNT
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Best Local
65338 CCACAGAGCTGAGGTAGAAGAAGGTATCAGAGAAGGGCAGGAGGATCATGTATGCCCTGA 65279
                                                                                                                                                                                                   65458 TGAAGCGGGCACGCTGGCGTTTCTCTTGGTTGGCATGCTGCAGAGTCAGGCGGCAGCAGA 65399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  misc_feature
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                                                                                                                                141 GCACCTGCACGAACACCCGCCGAAACTGCTGCGGGGGACACCCGTGTACAGGAGCGGGGTTGA 200
                                                                                                                                                                                                                                                              81
                                                                                                                                                                                                                                                                                                                                                      21 GGCGCCGGGACGCGAAGAGCAACGGGCGCTGCCACAAAGCGGGCGCTGTCGGTGGTGGAGT 80
                                                TGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGA 260
                                                                                                                                                                                                                                                                                                      GCCGGGAGGCTAGGAAGATGAGGGGGGCTGCGGCTGAGCTGGTGCTGTCCTTGGTGGAGA 65459
                                                                                                GCACCTGCCAGAACACCTTCCGGAACTGCTGAGAGGACACGTTGTAGAGGAGAGGGTTGA
                                                                                                                                                                                                                                                      GCGCATGTACGCGCAGGCGCTTCTCGTGGTGGTGGCGTGCTGCAGCGACAGGCGGCAGCACA 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chemistry: Dye-primer ET; 0% of reads Chemistry: Dye-terminator Big Dye; 100% of reads Assembly program: Phrap; version 0.99319 Consensus quality: 208484 bases at least 040 Consensus quality: 208911 bases at least 030 Consensus quality: 208985 bases at least 020
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Quality coverage: 0.00 in Q20 bases; agarose-fp Quality coverage: 11.62 in Q20 bases; sum of contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequencing vector: M13; 0%
                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Insert size: 206000; agarose-fp
Insert size: 212112; sum-of-contigs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequencing vector: plasmid; 100%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Center project name: M_BA0462P13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  59789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               arbitrary. Gaps between the configs are represented as runs of N. but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NOTE: This is a 'working draft' sequence. It currently consists of 5 contigs. The true order of the pieces is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11332
11432
28618
28718
49455
49555
107385
107385
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/note="assembly_name:Contig12"
.44784 c 46164 g 58745 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="assembly_name:Contig10"
49555. .107384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note="assembly_name:Contig11"
107485. .209885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="assembly_name:Contig8"
11432. .28617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Mus musculus"
/db_xref="taxon:10090"
/chromosome="UNK"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="assembly_name:Contig9"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone="RP23-462P13"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11331: contig of 11331 bp in length
11431: gap of unknown length
28617: contig of 17186 bp in length
28717: gap of unknown length
49454: contig of 20737 bp in length
49554: gap of unknown length
107384: contig of 57830 bp in length
107484: gap of unknown length
107486: gap of unknown length
209885: contig of 102401 bp in length
                                                                                                                                                                                                                                                                                                                                                                                                                                         48.3%;
76.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 302; DB 2; Length 209885; Pred. No. 9.4e-59;
                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches 131; Indels 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          403 others
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RESULT 13
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              Allen, C., Addons, C., Addons, C., Adlon, C., Are, J.R., Ayele, M., Banks, T., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbaria, J., Benton, J., Blanage, K., Blankenburg, K., Bonin, D., Bouck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Bouck, J., Bowie, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Bouck, J., Bowie, S., Brieva, M., Brown, E., Bryant, N.P., Bouck, J., Bowie, S., Brown, E., Bryant, N.P., Bouck, J., Chares, J., Chares, J., Chares, J., Chares, D., Chen, G., Chen, Z., Chowdhry, I., Christopoulos, C., Carron, T.F., Carter, M., Cavazos, S.R., Chacko, J., Chavez, D., Chen, G., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox, C., Davy-Carroll, J., Dederich, D.A., Davila, M., L., Davis, C., Davis, C., David, R., Davila, M., L., Ding, Y., Dinh, H.H., Duthwaite, K. J., Daper, H., Dugan-Rocha, S., Durbin, K. J., Earnhart, C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Falls, T., Ferraguto, D., Flagg, M., Ford, J., Foster, P., Frantz, P., Falls, T., Ferraguto, D., Flagg, M., Ford, J., Foster, P., Frantz, P., Garran, D., Dathwaite, K., J., Garcia, A., Garner, T., Garza, N., Gill, R., Gorrell, J.H., Garza, N., Gill, R., Gorrell, J.H., Garza, N., Gill, R., Gorrell, J.H., Garza, N., Gill, R., Gorrell, J.H., Hodgson, A., Hogues, M., Holloway, C., Hollins, B., Homsi, F., Howard, S., Huber, J., Hulyk, S., Hume, J., Jackson, L.E., Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S., Kratovic, J., Kureshi, A., Landry, N., Leal, B., Lewis, L.C., Lewis, L., Li, J., Liu, W., Loulseged, H., Lozado, R., J., Lucher, S., Liu, W., Loulseged, H., Lozado, R., Martin, R., Martinez, E., Mastinez, E., McLeod, M.P., Meador, M., Martinez, E., McLeod, M.P., Meador, M., Martinez, E., McLeod, M.P., Meador, M., Muller, S., Martinez, E., McLeod, M.P., Meador, M., Morris, S., Mosery, G., Wettse, A., Palon, R., Paten, Y., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Stone, H., Sto
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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Usmani,K., Vasquez,L., Vera,V., Villalon,D., Vinson,R., Wang,Q.,
Wang,S., Ward-Moore,S., Warren,R., Mashington,C., Watlington,S.,
Williams,G., Williamson,A., Wleczyk,R., Wooden,S., Worley,K.,
Wu,C., Wu,Y., Wu,Y.F., Zhou,J., Zorrilla,S., Nelson,D.,
Weinstock,G. and Gibbs,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (17-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA On Jul 13, 2002 this sequence version replaced g1:20303189.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of Molecular and Human
Baylor Plaza, Houston,
3 (bases 1 to 92874)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (19-FEB-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Worley, K.C.
Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished
                                                                                                                                                                                                                                                               arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence
                                                                                                                                                                                                                                                                                         (see http://www.hgsc.bcm.tmc.edu/docs/Genbank_draft_data.html).
NOTE: This is a 'working draft' sequence. It currently
consists of 50 contigs. The true order of the pieces
is not known and their order in this sequence record is
                                                                                                                                                                                                                                                                                                                                 NOTE: Estimated insert size may differ from sequence length
                                                                                                                                                                                                                                             as soon as it is available and the accession number will be preserved.
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                                                                                                                                                                                                                                                                                                                                              Sequencing vector: Plasmid;
Chemistry: Dye-terminator Big Dye: 100% of reads
Assembly program: Phrap; version 0.990329
Consensus quality: 32512 bases at least Q40
Consensus quality: 34515 bases at least Q30
Consensus quality: 35930 bases at least Q20
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Web site: http://www.hgsc.bcm.tmc.e
Submission
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Mus musculus brain cDNA, clone MNCD-VV/1.

AB041649.1 GI:7670499
fis (full insert sequence).
Mus musculus (strain:C57BL) adult female cDNA to mRNA,
Mus musculus (strain:C57BL) adult female cDNA to mRNA,
     Hashimoto, K.,
                                                                 Hashimoto, K. isolation of full-length cDNA
                                                                                                                                       Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
                                    Unpublished
                                                    made by oligo-capping
                                                                                                     Osada, N., Kusuda, J.,
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/db_xref="taxon:10116"
/clone="CH230-286021"
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     Kusuda,J.
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                 GGAGAGACAGAGAAAAGAAAAAAAA 433
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TGGGCATCCAACACGGCCAACGTCACCACAATCAGTCCT-GGCAGGCAAGAACAGGAG
                                                                                                  TGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAGGAG
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                                                                                                                                                          AGTAGGACCTCGTCCAGTCGTGCTTTGGGTTTTGGCCGCAGCCATGATCCTCCGAATCTGGT 320
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               312;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (13-APR-2000) Katsuyuki Hashimoto, National Institute of Infectious Diseases, Division of Genetic Resources; 23-1, Toyama 1-chome, Shinjuku-ku, Tokyo 162-8640, Javana 1-chome, Shinjuku-ku, Tokyo 162-8640, Javana (E-mail:khashi@nih.go.jp, URL:http://www.nih.go.jp/yoken/genebank/, Tel:81-3-5285-1111(ex.2120), Fax:81-3-5285-1181)
URL: http://www.nih.go.jp/yoken/genebank/
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//PICTEIN_Id="BAA95101.1"
//Gb_xref="GI:7670500"
/translation="MWVLGIAATFCGLFWLPGLALQIQCYQCEEFQLNNDCSSPEFIV
/CTVNVQDMCQKEVTEQSAGIMYRKSCASSAACLIASAGYQSFCSPGKLNSVCISCCN
TPLCNGPRPKKRGSSASAIRPGLLTTLLFFHLALCLAHC"
a 487 c 511 g 408 t
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/dev_stage="adult"
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/clone="MNCb-0671"
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DNA

NA linear HTG 19-JUL-2002
\*\*\* SEQUENCING IN PROGRESS

KEYWORDS VERSION ACCESSION DEFINITION Locus

ORGANISM

Rattus norvegicus Rattus norvegicus. HTG; HTGS\_PHASE1 AC128363.1

REFERENCE AUTHORS

Rattus

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Mazny, D.M., Mains, C., Adio-Oduola, B., Ali-osman, F.R., Allen, C., Alsbrooks, S.L., Amaratunge, H.C., Are, J.R., Ayele, M., Banks, T., Barbarla, J., Benton, J., Binage, K., Blankeaburg, K., Bonnin, D., Bertonk, J., Bonde, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Buck, J., Bowle, S., Brieva, M., Brown, E., Brown, M., Bryant, N.P., Buck, J., Bouck, J., Burch, P., Burkett, C., Burch, F., Chacko, J., Chavez, D., Chen, G., Chen, R., Chen, Z., Chowdhry, I., Christopoulos, C., Cleveland, C.D., Cox, C., Coyle, M.D., Dathorne, S.R., David, R., Delaney, K.R., Delgado, O., Denn, A.L., Ding, Y., Dinh, H.H., Douthwaite, K.J., Draper, H., Dugan-Rocha, S., Durbin, K.J., David, R., Delgado, O., Denn, A.L., Ding, Y., Dinh, K.J., David, R., C., Edgar, D., Edwards, C.C., Elhaj, C., Escotto, M., Falls, T., Ferraguto, D., Flagg, M., Ford, J., Foster, P., Frantz, P., Falls, T., Ferraguto, D., Flagg, M., Ford, J., Foster, P., Frantz, P., Gabisi, A., Gao, J., Garcia, A., Garner, T., Garza, N., Gill, R., Gorrell, J.H., Guevara, W., Gunaratne, P., Halles, S., Hamilton, K.J., Harris, K., Hart, M., Havlak, P., Hawes, A., Hernandez, J., Hernandez, O., Hodgson, A., Hogues, M., Holloway, C., Hollins, B., Homsi, F., Howard, S., Will, L., Korvah, J., Kovar, C., Karlsson, E., Klann, U., King, L., Kurdy, C., Hollins, B., Hemis, L.C., Levis, F., Howard, S., Hume, J., Jackson, L.E., Jacobson, B., Jia, Y., Johnson, R., Jolivet, S., Joudah, S., Karlsson, E., Machiney, E., Martin, R., Martindale, A., Martinez, E., Massey, E., Mawhiney, E., McLeer, A., Lucier, R., Luna, R., Martinez, E., Luna, R., Massey, E., Mawhiney, E., McLeer, A., Lucier, R., Luna, R., May, J., Perez, L., Peters, J., Pickens, R., Primus, E., Pull, L., Mohabbat, K., Moygen, M., Morris, S., Massey, E., Makhiney, E., MacLeer, A., Payton, B., Peery, J., Perez, L., Peters, J., Peters, J., Peters, J., Peters, J., Peters, J., Peters, J., Peters, J., Peters, J., Peters, J., Peters, J., Peters, J., Peters, J., Peters, J., Peters, J., Peters, J., Peters, J., Peters, J., Peters,
                                                                                                                                                    Submitted (19-JUL-2002) Human Genome Sequencing Center, Department of Molecular and Human Genetics, Baylor College of Medicine, One Baylor Plaza, Houston, TX 77030, USA
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**** 69 unordered pieces.
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                                                                                                                                                                                                                                                                (bases 1 to 190503)
 Center project Information
Center project name: GZCA
Center clone name: CH230-22808
Center Summary Statistics
                                                             Web site: http://www.hgsc.bcm.tmc.edu/
Contact: hgsc-help@bcm.tmc.edu
------Project Information
                                                                                                                                   Center: Baylor College of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              arbitrary. Gaps between the contigs are represented as runs of N, but the exact sizes of the gaps are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved.
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NOTE: This is a 'working draft' sequence. It currently consists of 69 contigs. The true order of the pieces is not known and their order in this sequence record is
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JOURNAL REFERENCE AUTHORS TITLE

Unpublished 2 (bases 1

Direct Submission

JOURNAL

Direct Submission Worley, K.C. TITLE

COMMENT

Center code: BCM

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NOTE: Estimated insert size may differ from sequence length
                                                               Sequencing vector: Plasmid; Chemistry: Dye-terminator Big Dye: 100% of reads Assembly program: Phrap; version 0.990329 Consensus quality: 131119 bases at least Q40 Consensus quality: 136808 bases at least Q30 Consensus quality: 141839 bases at least Q30
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Db 169311 GAAGATGAGGGGGCTGTGTGCTGAGCTAGTGCTGTCCTTGGTGGAGCTGAAGTGTGAACG
                                                                                                                                                   Query Match
Best Local Similarity
Matches 233; Conserv
                                         154 CACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAG
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| 2        | an       | ч        | 7TM recep |          |          |          | DNA encoding human | an full-l | DNA encoding human | G pro    |          | G-protein coupled | G protein-coupled | Human full-length | DNA encoding novel | Non-endogenous hum |          | Human ovarian canc | 0        | ovarian  | ovarian  | ovarian  | ovarian can | Human G protein co | noma     |          |          | Human ovarian canc |          | ovarian ca | Primer specific fo | cDNA ence | secreted |          | Human ovarian anti |

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## ALIGNMENTS

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WPI;

2002-500186/53.

Novel ovarian cancer polypeptide and polynucleotide, useful for

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence of the invention.
                     Ovarian
                                                                                ABL40348 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian cancer
                                         28-JUN-2002
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                     carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                       (first entry)
cancer therapy; vaccine;
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 gene therapy; tumour; cancer;
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The invention relates to ovarian tumour polynucleotides and polypeptides CC that may be utilised in cancer therapy, for example in a vaccine or CC gene therapy. Polypeptides and polynucleotides of the invention are CC expanding T-cells specific for a tumour protein, and for inhibiting the CC development of a cancer in a patient. They are also useful for Stimulating an immune response in a patient, and for treating a cancer in a patient and for determining the presence of a cancer in a patient.

CC The isolated polynucleotides of the invention are useful for their CC ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for dealyning and CC preparing ribozyme molecules for inhibiting expression of tumour cells. Polypeptides and polynucleotides of the CC invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current consumers of the ovarian carcinoma 01034C EST clone nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-SEP-1999;
01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
14-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                            (XUJJ/)
(STOL/)
(ALGA/)
                                                                                                                                                                                                                                                                                                                          Ovarian tumour polypeptide a prevention and/or treatment
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                                                                                                                                                                                                                                                                                                Claim
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) ALGATE P A
) FLING S P.
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2000US-0561778.
2000US-0640173.
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                                                                                                                                                                                                                                                                                                                          polypeptide and 
/or treatment of
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                                                                                                                                                                                                                                                                                                English.
                                                                                                                                                                                                                                                                                                                                                                                                   Fling
                                                                                                                                                                                                                                                                                                                          polynucleotide useful in diagnosis, cancer, especially ovarian cancer
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Sequence 625 BP; 182 A; 144 C; 182 G; 116 T; 1 other;

δÃ 멍 δÃ DЬ δõ В Q В Qy Matches Query Match Best Local Similarity 241 181 181 121 121 61 61 ш GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG CAGCGACAGGCGCAGCACCAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC CAGCGACAGGCGCAGCACCAGCACCTGCACGAACCCCGCCGAAACTGCTGCGAGGACAC GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG GCCCTGTCCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCCTGCACAAAGCG CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCCGCTGCACAAAGCG 625; Conservative 99.8%; 0; Score 624; Pred. No. Mismatches 4 DB .6e-154; 0, Length 0; Gaps 240 180 120 60 300 300 180 120 0,

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              The present invention describes a composition (I) comprising: carriers and immunostimulants; and a polypoptide (II) of a ovarian tumour polypoptide encoded by a polypuctide (III) having a CDNA sequence (S1) from the 10912 nucleotide sequences as given in ABL77023 to ABL87934, (III) encoding (II) having a sequence (S2), a T cell population of (II), or antigen presenting cells that express (II). (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to (S1) can be used for detecting ovarian cancer in a patient's biological sample preferably serum or ovarian tissue. The method comprises contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to
                                                                                                                                                                                                                                                                                                                                                                                                                                       polypeptide of a ovari
polypeptide, antibody
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                                                                                                                                                                                                                                                                                                                                                                Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                 for therapy and diagnosis of ovarian cancer comprising of a ovarian tumor polypeptide, polynucleotide encoding , antibody specific to polypeptide or T cell expressing
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05-SEP-2002

(first entry)

ABT03284 standard;

CDNA; 1897

cytostatic;

Human; ovarian cancer;

ovarian

carcinoma;

gene

therapy;

immunotherapy;

Human ovarian carcinoma associated coding sequence SEQ ID NO:

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Matches 625
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                                                   AACACTGTTTCAAACCCGGGGGGGTTGGCTGTGTTAAAGAAAAGACCATTAAATGCTTTAG
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Pred. No. 4.6e-154;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian cancer
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03-APR-2001;
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TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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                                          TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
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2001US-0825294.
2001US-0970966.
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99.8%;
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7.4e-154;
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RESULT 5
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01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
                                                                                                  The invention relates to ovarian tumour polynucleotides and polypeptide that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their
         ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the ovarian carcinoma 01034C/0591S consensus
                                                                                                                                                                                                                                          Claim la;
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                                                                                                                                                                                                                                                                  Ovarian tumour polypeptide and prevention and/or treatment of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ovarian cancer; cancer therapy; vaccine;
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) STOLK J A.
) ALGATE P A.
) FLING S P.
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2000US-0640173.
2000US-0656668.
2000US-0713550.
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/product= "Ovarian
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RESULT 6
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        13-NOV-2001; 2001WO-US45395
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                                                             Homo sapiens.
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atic; gene;
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Pred. No. 7.4e-154;
                                                                                   carcinoma; gene therapy;
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03-APR-2001; 2001US-0825294
02-OCT-2001; 2001US-0970966
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim
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          CTGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACA
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GACAGTGTAAAAAAAAAAAAAAAAA
                                                  GTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA
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                                                                                GTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                   The present invention provides human ovarian cancer associated proteins and coding sequences. The sequences can be used in the diagnosis and treatment of ovarian cancers. The present sequence is a coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                             Novel ovarian cancer polypeptide and polynucleotide, useful for detecting the presence of ovarian cancer in a patient, and in pharmaceutical compositions, e.g. vaccines, for treating ovarian
                                           1232
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                                                                                                                                                                                                                                                                                                                                                                               Claim
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03-APR-2001;
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          CCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC
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 CCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC
                                                                                    CCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGA
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                                                                                                                                                                                                                                                                                                        1619 BP;
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2001US-0825294.
2001US-0970966.
                                                                                                                                                                                                                                                              Conservative
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99.7%;
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01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer a patient and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary street of the entire desired gene or gene fragments, and for designing and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (XUJJ/)
(STOL/)
(ALGA/)
(FLIN/)
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                                                                                                                                                                                                    The invention relates to ovarian tumour polynucleotides and polypeptides that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are
                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI;
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                                                                                                                                                                                                                                                                                                                Claim la;
                                                                                                                                                                                                                                                                                                                                                            Ovarian tumour polypeptide and prevention and/or treatment of
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) ALGATE P A.
) FLING S P.
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2000US-0640173.
2000US-0656668.
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                                                                                                                                                                                                                                                                                                                                                            polynucleotide useful in cancer, especially ovaria
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ovarian cancer
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Best Local
           Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynuclectides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence represents the extended CDNA sequence of ovarian carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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                                                                                                                                                                                 GACAGTGTAAAAAAAAAAAAAAAAA 1617
                                                                                                                                                                                              CCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAA
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                                         cancer;
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57887 given
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                                                                                  (first entry)
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                                      cancer therapy; vaccine; gene therapy; tumour;
                                                                                                                             CDNA; 1619
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99.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      that may be utilised in cancer therapy, for example in a vaccine or gene therapy. Polypeptides and polynucleotides of the invention are useful for detecting a cancer in a patient, for stimulating and/or expanding T-cells specific for a tumour protein, and for inhibiting the development of a cancer in a patient. They are also useful for stimulating an immune response in a patient, and for treating a cancer in a patient, and for determining the presence of a cancer in a patient. The isolated polynucleotides of the invention are useful for their ability to selectively form duplex molecules with complementary stretches of the entire desired gene or gene fragments, and for designing and preparing ribozyme molecules for inhibiting expression of tumour polypeptides in tumour cells. Polypeptides and polynucleotides of the invention are also useful in recombinant DNA molecules to direct expression of a polypeptide in appropriate host cells. The current sequence or presents the content of the current sequence.
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01-MAY-2000;
15-AUG-2000;
07-SEP-2000;
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CCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCC
                                                        GGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCCGCAG
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                                                                                                                               CCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGA
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                                                                                                                                                                                                                                   GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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) STOLK J A.
) ALGATE P A.
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                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                            represents the ovarian carcinoma O591S nucleotide
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99.7%;
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                                                                                                                                                                                                                                                                                                                                                                     DB 24;
.4e-151;
                                                                                                                                                                                                                                                                                                                                                                                                                other;
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            The invention relates to 2175 novel human ovarian antigens (ABP41054-ABP43228) and to cDNAs encoding them (ABP4131-ABQ56305), and also encompasses polypeptides 90% identical and polynucleotides 95% identical to the sequences of the invention. The invention additionally relates to recombinant vectors and host cells comprising human ovarian antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABQ54231 standard;
                                                                                                       Claim
                                                                                                                                      useful in the prevention, treatment
ovarian cancer), immune disorders,
neurological diseases -
                                                                                                                                                 Isolated nucleic acid molecules encoding novel ovarian polypeptides, useful in the prevention, treatment and diagnosis of cancer (e.g. ovarian cancer), immune disorders, cardiovascular disorders and
                                                                                                                                                                                                                                                                                                              07-JUN-2000; 2000US-209467P
                                                                                                                                                                                                                                                                                                                                           07-JUN-2001; 2001WO-US18569
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                                                                                                                                 neurological
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                                                                                                    2922pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            reproductive; chromosome 2q21-22;
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   ovarian
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ischaemia, angiogenesis, nervous system disorders e.g. Alzheimer's disease, infections caused by bacteria, viruses and fungi and ocular disorders e.g. corneal infection. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin aging due to sunburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities. AAF22364 to AAF22372 and AAB63133 represent sequences used in the exemplification of
                                                                                                                                                                                                              cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; and ophthalmological. The polynucleotides and proteins can be used to prevent, treat or amelicrate a medical condition in e.g. humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. They are also used in diagnosing a pathological condition or susceptibility to a pathological condition. Disorders which are diagnosed or treated include autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; ophthalmological; gene therapy; autoimmune disease; neoplasm; rheumatoid arthritis; hyperproliferative disorder; cardiac arrest; cardiovascular disorder; cerebrovascular disorder; cerebral ischaemia; angiogenesis; nervous system disorder; Alzheimer's disease; infection; angiogenesis; nervous system disorder; Alzheimer's disease; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                         and polypeptides homologous to them. Human secreted proteins have activities based on the tissues and cells the genes are expressed in Examples of activities include: immunosuppressive; antiarthritic;
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20-JAN-2000; 2000US-0176931.
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                                                                                                                                                                 Human; secreted protein; immunosuppressive; antiarthritic; antirheumatic; antiproliferative; cytostatic; cardiant; vasotropic; cerebroprotective; nootropic; neuroprotective; antibacterial; virucide; fungicide; opthalmalogical; vulnerary; autoimmune disease; cardiovascular disorder; hyperproliferative disorders; cerebrovascular disorder; wound healing; hyperproliferative disorders; cerebrovascular disorder; wound healing;
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99.78;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC vasottopic; cerebroprotective; nontropic; neuroprotective; antibacterial; cc vasottopic; cerebroprotective; nontropic; neuroprotective; antibacterial; cc virucide; fungicide; opthalmalogical; and vulnerary. The proteins, cc polynucleotides, agonists and antagonists can be used to treat or detect or diagnose various diseases and disorders including, autoimmune cc diseases e.g. rheumatoid arthritis, hyperproliferative disorders cc e.g. neoplasms of the breast or liver, cardiovascular disorders cc e.g. cardiac arrest, cerebrovascular disorders e.g. Alzhelmer's disease, cc infections caused by bacteria, viruses and fungi and ocular disorders c.g. corneal infection. The polypeptides can also be used to aid wound cc healing and epithelial cell proliferation, to prevent skin ageing due to sunburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities. Included in the invention are polynucleotide sequences AaF64167 - AaF64175 and peptide AaB75505 which cc are used in the isolation, identification and characterisation of the
                                                                                                                                                                                                                                                                                                                                                                                        Matches 606;
                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human secreted proteins AAB75506 - AAB75554 are encoded by polynucleotid sequences AAF64176 - AAF64224. The specification includes amino acid sequences AAB75555 - AAB75606 which represent fragments of the human secreted proteins, and protein sequences with which they share homology.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acid molecules encoding human secreted proteins, preventing, treating or ameliorating a disorder, e.g. Al Parkinson's diseases and cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expressed, examples of these activities include, immunosuppressive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The proteins and polynucleotides, their agonists and antagonists have activities dependent on the tissues and cells in which they are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1956 BP; 430 A; 543 C; 524 G; 456 T; 3 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antiarthritic; antirheumatic; antiproliferative; cytostatic;
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                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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AAB88317 - AAB88419. Included in the invention are primers AAF93917 - AAF94295 and AAF62235 - AAF62235 which are used to isolate to CDNA sequences of the invention. The invention also includes methods for the production of antibodies directed against the proteins, and cDNA sequences, which can be used in vaccines. The polynucleotide sequences can be used in gene therapy. The polynucleotide sequences and the proteins they encode may be used in the prevention, treatment and diagnosis of diseases associated with inappropriate secretory protein/membrane protein expression. The nucleic acids and complementar
                                                                                                             This invention relates to nucleic acid sequences AAF93744 - Awhich encode human secretory or membrane proteins represented
                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; secretory protein; membrane protein; vaccine; gene therapy; rheumatoid arthritis; diabetes; ss.
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11-JAN-2000;
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   complementary
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Best Local
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Human; secretory protein; membrane protein;
rheumatoid arthritis; diabetes; PCR primer;
                                          Primer specific for DNA encoding secretory/membrane
                                                                                                                                           AAF94186 standard;
                                                                                                                                                                                                                        1830
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (e.g. polymerase chain reactions (PCR)) to detect and quantitate the presence of similar nucleic acid sequences in samples. They may also be used to study the expression and function of secretory proteins/membrane polypeptides and their role in metabolism. The polypeptides may be used as antigens in the production of antibodies against them and in assays to identify modulators (agonists and antagonists) of expression and activity. The antibodies and antagonists may also be used as therapeutic agents to down regulate expression and activity. The antibodies may also be used as diagnostic agents for detecting the presence of the polypeptides in samples (e.g. by enzyme linked immunosorbant assay (ELISA). Examples of diseases which may be treated include rheumatoid arithritis and diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1590
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1890 BP; 419 A; 528 C; 533 G; 410 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  arthritis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequences may also be used as DNA probes in diagnostic assays (e.g. polymerase chain reactions (PCR)) to detect and quantit
                                                                                                                                                                                                                                                                                                                                                                               427
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7 CCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCGGGCGCT 66
                                                                                                                                                                                                                CCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCCTGGGCA 366
                                                                                                                                                                                                                                                                                                          GACACGAGCAGGAGAGAGAGAAAAAGAAAAACACAGCATGAGAACACAGTAAATA 1709
                                                                                                                                                                                                                                                                                                                                                                                                                        CATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGAT 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CAGGCGGCAGCACCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTA 186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGA 126
                                                                                                                                                                                                                                                                                                                                                             AATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACC 486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCCGAGAAGGGGAGGAGGAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CAGGCGGCAGCACCAGCACCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTA 1469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGA 1409
                                                                                                                                                                                                                                                                                AATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       599;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diabetes.
                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                95.7%;
99.8%;
                                                                             entry)
                                                                                                                                           587
                                                                                                                                           ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 598.4; DB 22; Pred. No. 3.9e-147;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
             vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 1890;
                                            protein SEQ
             gene therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                           cc which encode human secretory or membrane proteins represented by
cc AAR93917 - AAR94295 and AAR62232 - AAR62235 which are used to isolate the
cd AAR93917 - AAR94295 and AAR62232 - AAR62235 which are primers
cc AAR93917 - AAR94295 and AAR62232 - AAR62235 which are used to isolate the
cd AAR93917 - AAR94295 and AAR62232 - AAR62235 which are used to isolate the
cc cDNA sequences of the invention. The invention also includes methods for
cc sequences, which can be used in reactions. The polynucleotide sequences
cc can be used in gene therapy. The polynucleotide sequences and the
cc diagnosis of diseases associated with inappropriate secretory
cc proteins they encode may be used in the prevention, treatment and
cc diagnosis of diseases associated with inappropriate secretory
cc sequences may also be used as DNA probes in diagnostic assays
c (e.g. polymerase chain reactions (PCR)) to detect and quantitate the
cc used to study the expression and function of secretory proteins/membrane
cc used to study the expression and function of secretory proteins/membrane
cc used to study the expression and function of secretory proteins/membrane
cc identify modulators (agonists and antagonists) of expression and
cc agents to down regulate expression and activity. The antibodies may also
be used as diagnostic agents for detecting the presence of the
cc polypeptides in samples (e.g. by enzyme linked immunosorbant assay
cc (ELISA). Examples of diseases which may be treated include rheumatoid
carthritic and diseases which may be treated include rheumatoid
                                                                                                                                                                                                                                                                                                                                                Matches 498;
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 587 BP; 117 A; 162 C; 140 G; 162 T; 6 other;
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11-JAN-2000; 2000JP-0118775
02-MAY-2000; 2000JP-0183766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acids encoding secretory proteins/membrane pro
       330
                                                                                                                                                                                                                                                                         516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention relates to nucleic acid sequences AAF93744 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 5; SEQ ID 620; 609pp + CD ROM; English
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                                             338
                                                                                   270
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                                                                                                                                                                                                                                   150
                                                                                                                                                                                                                                                                                                       90 CGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGGCAGCAGCACCAGCACCTGCA 149
                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                 TCGTCCAGTCGTGCTTGGGCTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCC
AGCATACGGCCAATGTCACAACAATCAGCCCTGGGCAGACACGAGCAGGAGGAGAGACA
                                                                                                              TGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAAGGATCATGTACGCCCGGAAGTAGGACC
                                                                                                                              TGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACC
                                                                                                                                                                                                             CGAACACCCGCCGAAACTGCTGCGAGGACCACCGTGTACAGGAGCGGGTTGATGACCGAGC
                                                                                                                                                                                                                                                                 CGCNCAGGCGCTTCTCGTGGTTGGCGTGTTCAAGCGACAGCGGCAGCC--AGCACNTGCA
                                                                                                                                                                                        CGACCACCCGCCGAANTTGTTGCGAGGACACCGTGTACAGAAGCGGGTTGATGACCGAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Isogai T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               and diabetes.
                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                76.78;
96.38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nishikawa
                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                            Score 479.4; DB 22; Length 587;
Pred. No. 4.4e-116;
0; Mismatches 17; Indels 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          proteins, useful
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The present invention describes a composition (I) comprising: carriers can dimmunostimulants; and a polypeptide (II) having a cDNA sequence CC polypeptide encoded by a polypuciectide (III) having a cDNA sequence CC (S1) from the 10912 nucleotide sequences as given in ABI/7023 to CC ABL87934, (III) encoding (II) having a sequence (S2), a T cell CC population of (II), or antigen presenting cells that express (II). CC (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to CC (S1) can be used for detecting ovarian cancer in a patient's biological Sample preferably serum or ovarian tissue. The method comprises CC contacting a biological sample from a patient with (IV), detecting the amount of polynucleotide hybridising to (IV) and comparing the amount to a predetermined cutoff value and thereby detecting ovarian cancer in the CC patient, where the amount of polynucleotide hybridising to (IV) is CC detected preferably by polymerase chain reaction (PCR). (I) comprising CC (III) and/or (II) is useful for stimulating and/or expanding T cells specific for an ovarian tumour protein comprising contacting T cells with (III) or (II), is useful in design and preparation of cribozyme molecules for inhibiting expression of the tumour polypeptides and proteins in tumour cells; and to isolate a full length gene from a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABL79397 standard; cDNA; 558 BP
                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID 2375; 489pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-122075/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Algate PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-MAY-2001; 2001WO-US17756
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; ovarian cancer; ovarian tumour; cytostatic; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human ovarian cancer related cDNA clone SEQ ID NO:2375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-MAY-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                polypeptide
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Best Local
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533 NTTAACATTGTTTCAANCCGGGGG
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                                                        CAGTAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGA
                                                                                                                                          CCTGGGCAGACACGAGCAGGAANGGAGAGAGAGAAAAAGAAAAACACAGCATGAGAACA
                                                                                                                                                                                        CATTGATTCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGC
                                                                                                                                                                                                     CCATGA-TCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGC
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                     TTTAACACTGTTTCAAACCCGGGG
                                             AATGGTACC-ATTTTTCAGTGTTTGACTTNACAGCTTCTTTTGCCACAAGCAAGAGAG-A
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Pred. No. 5.
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Search completed: November Job time: 135.06 secs 7, 2002, 10:10:19

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Title:
Perfect score:
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Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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625
24791104 seqs, 12571243825 residues
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Pending_Patents_NA_Main: *
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Copyright (c) 1993 - 2002 Compugen Ltd
                                                                                                                                                                                                                                                                           /cgn2_6/ptodata/1/pna/US082_COMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

0 0 0 0 0 0 0 0 0 0 0

997.3 997.3 997.1 997.1 997.1 997.1 997.1 997.1 998.6 998.6 998.6 998.6 998.6 998.6

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APPLICANT: COTIXA COTPORATION

APPLICANT: Xu, Jiangchun

APPLICANT: Stolk, John A.

APPLICANT: Stolk, John A.

APPLICANT: Filing, Steven P.

APPLICANT: Molesh, David Alan

APPLICANT: Molesh, David Alan

APPLICANT: Molesh, David Alan

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THE

TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER

FILE REFERENCE: 210121.48401PC

CURRENT APPLICATION NUMBER: PCT/US01/45395

CURRENT FILING DATE: 2001-11-13

NUMBER OF SEQ ID NOS: 215

SOFTMARE: FastSEQ for Windows Version 4.0

SEQ ID NO 210

LENGTH: 625

TYPE: DNA

ORGANISM: Homo sapiens
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                                                                                                    CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAAG
                                                                                                                                    GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 120
                                                                                                                                                                                                                                          99.8%; Score 624; DB 1; ilarity 100.0%; Pred. No. 5e-116; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 70 US-60-261-974-9
PCT-US01-18569-111
13 US-10-264-049-111
25 US-09-652-121-7293
25 US-09-652-128-9375
26 US-09-652-1108-9375
27 US-09-659-997-11085
28 US-09-710-281-4458
29 US-09-710-281-4458
20 US-09-726-805-1608
20 US-09-726-805-1914
20 US-09-950-083-1914
20 US-09-950-083-1914
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6 US-09-950-083-1915

US-10-155-299-2415

2 US-10-208-408-26

3 US-60-295-262-26

4 US-60-308-868-26

7 US-9371-168-7188

US-99-644-873-9077

US-99-652-109-9370
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US-09-359-922-11555
US-09-919-002-11555
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Sequence 11555, A
Sequence 1111, App
Sequence 1111, App
Sequence 1111, App
Sequence 1111, App
Sequence 3345, Ap
Sequence 3458, Ap
Sequence 4458, Ap
Sequence 1608, App
Sequence 1914, Ap
Sequence 1914, Ap
Sequence 2414, Ap
Sequence 2514, Ap
Sequence 26, App1
Sequence 26, App1
Sequence 26, App1
Sequence 2788, App
Sequence 2788, App1
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Sequence 2788, App1
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Sequence 2788, App1
Sequence 2788, App1
Sequence 2778, Ap
Sequence 3977, Ap
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                                                                                                    120
                                                                                                                                                                        60
                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(625)
OTHER INFORMATION: n = A
US-09-825-294-210
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                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CF
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT APPLICATION NUMBER: 215
SOTTWARE: FastSEQ for Windows Version 3.0
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Matches 625
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LENGTH: 625
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                              CAGCGACAGGCGGCAGCAGCACCTGCACGAACACCCGGCGGAAACTGCTGCGAGGACAC
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ilarity 100.0%;
Conservative (
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;; Pred. No. 5e-116;
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RESULT 1
PCT-US01-45395-210
Sequence 210, Application PC/TUS0145395; GENERAL INFORMATION:

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181

121 61 61 ; FEATURE: ; NAME/KEY: misc\_feature ; LOCATION: 607 ; OTHER INFORMATION: n = PCT-US01-45395-210

A,T,C

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Query Match
Best Local Similarity
Matches 625; Conser

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                                                                                                                                                                                  ; NAME/KEY: misc_feature
; LOCATION: 607
; OTHER INFORMATION: n =
US-09-867-701-10876
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                                                                                                                                     Query Match
Best Local Sim
Matches 625;
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GENERAL INFORMATION:
APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
TITLE REFERENCE: 210111.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 10876
LENGTH: 625
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
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AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG
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                     CAGCGACAGGCGGCAGCACCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACAC 180
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                                                                                                                                     Conservative
                                                                                                                                                            99.8%;
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                                                                                                                                    Score 624; DB 33; ; Pred. No. 5e-116; 0; Mismatches 0;
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                                                                                                                                                         Length 625;
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US-09-970-966-210
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Best Local S
Matches 625
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CURRENT FILING DATE: 2001-10-02
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 210
LENGTH: 625
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Stolk, John A.
APPLICANT: Molesh, David Alan
APPLICANT: Molesh, David Alan
APPLICANT: Fling, Steven P.
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C6
                                                                                                                                                                                                                                 ORGANISM: Homo sapiens FEATURE: NAME/KEY: misc_feature LOCATION: 607
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                                                                                                                                                      Score 624; DB 36;
; Pred. No. 5e-116;
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PCT-US01-45395-214
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GENERAL INFORMATION:
APPLICANT: Corixa Corporation
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 214
LENGTH: 1897
                                                                                                                                                                                                                                                                          APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
APPLICANT: Molesh, David Alan
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.48401PC
CURRENT APPLICATION NUMBER: PCT/US01/45395
CURRENT FILING DATE: 2001-11-13
NUMBER OF SEQ ID NOS: 215
                                                                                                                                                              Query Match
Best Local
                                                                                                                                                 Matches 624;
                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapiens
                                                                                            1271
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                                        GGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG
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Similarity
                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                       for Windows Version 4.0
                                                                                                                                                             99.8%;
                                                                                                                                               Score 624; DB 1;
Pred. No. 6.3e-116;
0; Mismatches 1;
                                                                                                                                                                        Length 1897;
                                                                                                                                                 Indels
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| Qy                                                              | Qy<br>Db                                                         | Db Qy                                                            | Quer<br>Best<br>Matc                                                                                                                     | CURR<br>CURR<br>NUMN<br>SEQ<br>SEQ<br>TY<br>TY<br>TEE<br>OR<br>NAME<br>OFFE<br>LC<br>OFFE<br>TO<br>TO<br>TO<br>TO<br>TO<br>TO<br>TO<br>TO<br>TO<br>TO<br>TO<br>TO<br>TO                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        | GENE APP APP APP APP APP APP TIT                                                                                                                                                                                                                           | RESULT<br>US-10-                           | дb | o p                                                                                         | Qy Y | В                                                   | ΩУ  | φ<br><b>Q</b>                                                    | Db                                                                                                                                                                                                 | Оу                                     | D 43                                                           | D 2                                                            |     | P &                                                              |
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| CAGCGACAGGCGCAGCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC 180 | GCCGCTGTCGGTGGAGTGCGCATGTACCCGCAGCGCTTCTCGTGGTTGGCGTGCTG 120<br> | AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCCTGCACAAAGCG 60 | Match 99.8%; Score 624; DB 42; Length 625; ocal Similarity 100.0%; Pred. No. 5e-116; s 625; Conservative 0; Mismatches 0; Indels 0; Gaps | TILE REFERENCE: 210121.484C7 CURRENT APPLICATION NUMBER: US/10/212,677 CURRENT FILING DATE: 2002-08-02 NUMBER OF SEQ ID NOS: 288 SOFTWARE: FASTSEQ for Windows Version 4.0 SEQ ID NO 210 LENGTH: 625 TYPE: DNA ORGANISM: Homo sapiens FEATURE: NAME/KEY: misc_feature LOCATION: 607 OTHER INFORMATION: n = A,T,C or G US-10-212-677-210                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        | INFORMATION: ANT: Chenault, Ruth A. ANT: Chenault, Ruth A. ANT: Xu, Jiangchun ANT: Fanger, Gary R. ANT: Harlocker, Susan L. ANT: MCNEILL, Patricla D. DE INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY DE INVENTION: AND DIAGNOSIS OF OVARIAN CANCER | -677-210<br>= 210, Application US/10212677 |    | ACACTGITTCAAACCCGGGGGAGTTGGCTGTGTTAAAGAAAGACCATTAAATGCTTTAG 600 ACAGTGNAAAAAAAAAAAAAAAA 625 | Ò    | GGTACCAATTTTTCAGTGTTGGACTTGACAGCTTCTTTTGCCACAAGCAAG | . G | TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480 | TGGGCAGACACGAGCAGGAGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAGAGCATGAGAACACAGCATGAGAACACAGCATGAGAACACAGAGAGAAAACACAGAGAGAAAAACACAGAGAGAAAAACACAGAGAGAAAAACACAGAGAGAAAAACACAGAGAGAAAAACACAGAGAGAAAAAA | GGGCAGACACGAGCAGGAGAGAGAGAGAGAAAAGAAAA | CATGATICCTCCGAATCTIGGECATCCAGCATACGGCCATGTCACCACCAATCAGCCC 300 | AGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC 30 | );  | CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG 240 |

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APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CF.
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 214
LENGTH: 1897
TYPE: DNA
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US-09-825-294-214
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Best Local Sim
Matches 624;
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 214
LENGTH: 1897
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APPLICANT: Fling, Steven P.
APPLICANT: X1, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.484C6
CURRENT APPLICATION NUMBER: US/09/970,966
CURRENT FILING DATE: 2001-10-02
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Pred. No. 6.3e-116;
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LENGTH: 1897
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: MCNeill, Patricia D.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
EILE REFERENCE: 210121.484C7
CURRENT APPLICATION NUMBER: US/10/212,677
CURRENT FILING DATE: 2002-08-02
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APPLICANT: Xu, Jiangchun
APPLICANT: Fanger, Gary F
APPLICANT: Harlocker, Sus
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Harlocker, Susan L.
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                                                                                                                                                                                                                                                               99.8%;
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                                                                                                                                                                                                                                                               Score 624; DB 42;
Pred. No. 6.3e-116;
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                                                                                         PRIOR APPLICATION NUMBER: US 60/323,739
PRIOR APPLICATION NUMBER: US 09/488,725
PRIOR PPLICATION NUMBER: US 09/488,725
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: US 09/552,317
PRIOR APPLICATION NUMBER: PCT/US00/35017
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: US 09/491,404
PRIOR APPLICATION NUMBER: US 09/491,404
PRIOR APPLICATION NUMBER: PCT/US01/02623
PRIOR APPLICATION NUMBER: PCT/US01/02623
PRIOR APPLICATION NUMBER: PCT/US01/02623
PRIOR FILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
PRIOR FILING DATE: 2001-01-25
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                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Novel Nucleic TITLE OF INVENTION: Polypeptides FILE REFERENCE: 809ACIP PCT
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                                                            APPLICATION NUMBER: PCT/US01/02623 FILING DATE: 2001-01-25 APPLICATION NUMBER: US 09/496,914 FILING DATE: 2000-02-03
FILING DATE: 2000-04-27
APPLICATION NUMBER: PCT/US01/03800
FILING DATE: 2001-02-05
                                               APPLICATION NUMBER: US
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Zhou, Ping
Zhou, Ryle V
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Zhao, Qing A.
Wang, Jian-Rui
xue, Aidong J.
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Drmanac, Radoje T
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Weng, Gezhi
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Ghosh, Malabika
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Best Local S
Matches 617
                                                                  Sequence 205, Application PC/TUS0145395 GENERAL INFORMATION:
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NUMBER OF SEQ ID NOS: 992
SOFTWARE: pt_FL_genes Version 6.0
SEQ ID NO 16
LENGTH: 1967
TYPE: DNA
                                            APPLICANT: Corixa Corporation APPLICANT: Xu, Jiangchun
  APPLICANT:
                      APPLICANT:
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PRIOR FILING DATE: 2000-02-28
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LOCATION: (338)..(763)
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Stolk, John A.
Algate, Paul A.
Fling, Steven P.
Molesh, David Alan
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Pred. No. 2.2e-114;
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; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
; TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
; FILE REFERENCE: 210121.48401PC
; CURRENT APPLICATION NUMBER: PCT/US01/45395
; CURRENT FILING DATE: 2001-11-13
; NUMBER OF SEQ ID NOS: 215
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 205
; SEQ ID NO 205
; TYPE: DNA
; ORGANISM: Homo Sapiens
PCT-US01-45395-205
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                                                 Sequence 211, Application PC/TUS0145395
GENERAL INFORMATION:
APPLICANT: Corixa Corporation
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
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Best Local Similarity
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                         APPLICANT:
 TITLE
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T: Fling, Steven P.
T: Molesh, David Alan
INVENTION: COMPOSITIONS
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Sequence 205, Application US/09713550

GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE OF INVENTION: THERAPY AND DIAGNOSIS OF OV.
FITLE REFERENCE: 210121.484C4
CURRENT APPLICATION UNUMBER: US/09/713,550
CURRENT FILING DATE: 2000-11-14
NUMBER OF SEQ ID NOS: 205
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US-09-713-550-205
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CURRENT APPLICATION UNMBER: PCT/US01/45395
CURRENT FILING DATE: 2001.11-13
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 211
LENGTH: 1619
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Best Local Similarity
Matches 624; Conserv
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Pred. No. 1e-113;
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APPLICANT: XU, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIA
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION UNMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 205
LENGTH: 1619
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APPLICANT: Xu, Jiangchun
APPLICANT: Stolk, John A.
APPLICANT: Algate, Paul A.
APPLICANT: Fling, Steven P.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF OVARIAN CAN
FILE REFERENCE: 210121.484C5
CURRENT APPLICATION NUMBER: US/09/825,294
CURRENT FILING DATE: 2001-04-03
NUMBER OF SEQ ID NOS: 215
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 211
LENGTH: 1619
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                        RESULT 15
US-09-825-294-211
Sequence 211, Application US/09825294
GENERAL INFORMATION:
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US-09-825-294-205
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| 31.6<br>31.6<br>31.6                                           | 31.6<br>6                                                   | 31.8<br>31.8                                               | 32.6<br>32.4<br>32.4<br>32.3                                                                      | 33333<br>3323<br><br>88888                                                                            |
| 5.11                                                           | 5555                                                        | 555                                                        | 5.5.5.5.<br>5.2222                                                                                | 55555                                                                                                 |
| 2845<br>2845<br>2845                                           | 2739<br>2080<br>2158                                        | 2689<br>308<br>831                                         | 1926<br>3096<br>3949<br>415<br>1335                                                               | 473<br>1024<br>1201<br>1242<br>1772                                                                   |
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|                                                                | 30,<br>127                                                  | Sequence 1, Appli<br>Sequence 18015, A                     |                                                                                                   | Sequence 14061, A<br>Sequence 1, Appli<br>Sequence 334, App<br>Sequence 5, Appli<br>Sequence 3, Appli |

## ALIGNMENTS

US-10-230-437-15

Sequence 15, Applica GENERAL INFORMATION: APPLICANT: Baker, K

Application US/10230437

APPLICANT:
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APPLICANT:
APPLICANT:

Goddard, Audrey

Godowski, Paul J. Grimaldi, J. Christopher Gurney, Austin L. Smith, Victoria Gerritsen, Mary

Kevin

APPLICANT:

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APPLICANN: Stephan, Jean-Philippe F.
APPLICANN: Watanabe, Colin L.
APPLICANN: Wood, William I.
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APPLICANN: WOOD, William I.
APPLICANN: WOOD, WILLIAM CATES AND MUGLEIC PILE REFERENCE: P330P1094
CURRENT FILIMG DATE: 2002-08-28
PRIOR PPLICATION NUMBER: 10/119,480
PRIOR APPLICATION NUMBER: 10/059113
PRIOR FILING DATE: 1997-09-17
PRIOR APPLICATION NUMBER: 60/062287
PRIOR FILING DATE: 1997-10-17
PRIOR PPLICATION NUMBER: 60/062287
PRIOR PPLICATION NUMBER: 60/062287
PRIOR PPLICATION NUMBER: 60/062887
PRIOR APPLICATION NUMBER: 60/062887
PRIOR APPLICATION NUMBER: 60/064103
PRIOR APPLICATION NUMBER: 60/068973
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/079294
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PRIOR PILING DATE: 1998-03-26
PRIOR PILING DATE: 1998-03-27
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Sequence 38, Application US/10270333
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APPLICANT: Cravchik, Anibal
TITLE OF INVENTION: ISOLATED G-PROTEIN COUPLED RECEPTORS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING GFOR PROTEINS, AND USES
TITLE OF INVENTION: THEREOF AS INSECTICIDAL TARGETS
FILE REFERENCE: CL000733CON
CURRENT FILING DATE: 2002-10-15
CURRENT FILING DATE: 2002-10-15
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                                                                                                                                                                                                                                                                                              ; TYPE: DNA
; ORGANISM: Drosophila
US-10-270-333-38
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SOFTWARE: FastSEQ for
SEQ ID NO 38
LENGTH: 2196
                                                                                                                                                                                                                        Matches 100;
                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/168 677
PRIOR FILING DATE: 1999-12-03
PRIOR APPLICATION NUMBER: 60/175,691
PRIOR FILING DATE: 2000-03-23
PRIOR FILING DATE: 2000-03-23
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                                                                                                                                              1888 CGCCTGCCGTTGGATGCTTGCCTCTGATGGTAGTGGTGGCGATGGTGGTGATGTGGCCCGTT 1829
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                      183 TGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGA 242
                                                                                                        123 GCGACAGGCGGCAGCACCACCCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 361 TGGGCAGACACGAGCAGGAGGAGAGACAGAGA
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                                                                                                                                                                  63 CGCTGTCGGTGGTGGGGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCA 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61
                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCGCTGCACAAAGCG
                                                                       GGCCCAGACAAGTGAGCAGCAGCCGCCAGAAGCCGCTGCGAAATTTGGAGGACATCAGGT 1769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CAGCGACAGGCGGCAGCACAGCACCTGCACGAACACCCGCCGAAAACTGCTGCGAGGACAC
TGTAGAGGATCGGGTTCATGGCTGAGTTTAGGTAGAGCATGAAGCGCGAGAAGTACAGCA 1709
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                                                                                                                                                                                                                                                                                                                                                                                    FastSEQ for Windows Version
                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                    7.48;
52.68;
                                                                                                                                                                                                                 0;
                                                                                                                                                                                                      Score 46; DB 6; Lengum Zarr. Pred. No. 0.0025; unamatches 90; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  393
                                                                                                                                                                                                                                                      Length 2196;
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                                                                                                                                                                                                                   Gaps
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US-60-416-172-67
Sequence 67, Application US/60416172
GENERAL INFORMATION:
APPLICANT: Chang, Hwai Wen
APPLICANT: Felts, Katherine A.
; OTHER INFORMATION: Human US-60-416-172-67
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US-10-270-333-37
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Sequence 37, Application US/10270333
GENERAL INFORMATION:
APPLICANT: Cravchik, Anibal
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CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/15,691
PRIOR APPLICATION NUMBER: 60/175,691
PRIOR FILING DATE: 2000-01-12
PRIOR FILING DATE: 2000-03-23
PRIOR FILING DATE: 2000-03-23
                                                                               SEQ ID NO 67
LENGTH: 667
TYPE: DNA
ORGANISM: Gallus gallus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 37
LENGTH: 6472
                                                                                                                                                             APPLICANT: Warren, Amy J.
APPLICANT: Cooper, Judith R.
APPLICANT: Cooper, Judith R.
TITLE OF INVENTION: Genes from Human Cancer Cells and
FILE REFERENCE: 216019-153
CURRENT APPLICATION NUMBER: US/60/416,172
CURRENT FILING DATE: 2002-10-04
NUMBER OF SEQ ID NOS: 141
SOFTWARE: PatentIn version 3.1
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Best Local Similarity
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TITLE OF INVENTION: THEREOF AS INSECTICIDAL TARGETS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: CL000733CON
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                                        NAME/KEY: misc_feature
                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               243 GGATCATGTA 252
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGTTGTAGTA 1699
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52.6%;
                     extended sequence
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                     for VOG3_71
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                                                                                                                                                                                                                                                                        Human
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Matches

Local Similarity

Conservative

0; Mismatches 160; Indels

0;

Gaps

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7.2%;

Score 45; DB 7; Pred. No. 0.0026;

DB 7; Length 667

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GENERAL INFORMATION:
APPLICANT: Chu, Peter
                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/284,763
PRIOR FILING DATE: 2001-04-18
PRIOR APPLICATION NUMBER: US 09/971,228
PRIOR FILING DATE: 2001-10-03
NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Zhao, Haoran
APPLICANT: Rigel Pharmaceuticals, Inc.
TITLE OF INVENTION: EDG: Modulators of Lymphocyte Activation and Migration
FILE REFERENCE: 021044-000310PC
CURRENT APPLICATION NUMBER: PCT/US02/12342
CURRENT FILING DATE: 2002-08-26
CURRENT FILING DATE: 2002-08-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                         NAME/KEY: CDS
LOCATION: (9)..(1031)
OTHER INFORMATION: mutant #1 human EDG1
                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                             OTHER INFORMATION:
                                                                                                                                                                                                                                              OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                   LENGTH: 1032
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          487 C 487
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121 CAGCGACAGGCGCAGCACAGCACCTGCACGAACACCCGGCGGAAACTGCTGCGAGGACAC 180
                                        Local Similarity 63.4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGT 345
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GGGTGAGAAGGACACCCGGCCCTGAGGCACAGGGACCTGCAGGGGACCCGACAGCGCTCGT 486
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Pardo, Jorge
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Spencer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Liao, X.
                                                                                                                                                                                                                         mutant #1, C-terminally truncated variant of
human endothelial differentiation G-protein
coupled receptor (GPCR) 1 (EDG1)
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                                                              6.2%;
                                          0
                                                              Score 38.6; DB Pred. No. 0.17;
                                          Mismatches
                                                                                 DB 1;
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                                                                                   Length 1032;
                                             Indels
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                                                                                                                                                                                                                                                                                     PCT-US02-12342-4/c
                                                                                                                                                                                                                                                                                                     RESULT 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
LENGTH: 1149
                                                                                                                                                                                                                                             Sequence 4, Application PC/TUS0212342 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT FILING DATE: 2002-08-26
PRIOR APPLICATION NUMBER: US 60/284,763
PRIOR FILING DATE: 2001-04-18
PRIOR APPLICATION NUMBER: US 09/971,228
PRIOR FILING DATE: 2001-10-03
NUMBER OF SEQ ID NOS: 16
                                                                                        APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: PCT/US02/12342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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                APPLICANT: Spencer, Colin
APPLICANT: Zhao, Haoran
APPLICANT: Rigel Pharmaceuticals, Inc.
APPLICANT: Rigel Pharmaceuticals, Inc.
TITLE OF INVENTION: EDG: Modulators of Lymphocyte Activation and Migration
FILE REFERENCE: 021044-000310PC
CURRENT APPLICATION NUMBER: PCT/US02/12342
                                                                                                                                                APPLICANT:
                                                                                                                                                                                      APPLICANT: Chu, Peter APPLICANT: Jiang, Yir APPLICANT: Li, Congfe
CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: wild-type human endothelial differentiation OTHER INFORMATION: G-protein coupled receptor (GPCR) 1 (EDG1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: CDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1004 CGGGCACTTGCAGCAGGACATGATCCGGATGAAGGCCCGACGCATCTCCTTGTTGGTCAG 945
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              121 CAGCGACAGGCGGCAGCACAGCACCTGCACGAACACCCGGCGGAAACTGCTGCGAGGACAC 180
                                                                                                                                                                                                                                                                                                                                                                 AGTGTAAATGATGGGGTTGGTGCCGGAGTTGAG
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Zhao, Haoran
Rigel Pharmaceuticals, Inc.
Rigel Pharmaceuticals, Inc.
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Rigel Pharmaceuticals, Inc.
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Pardo, Jorge
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                                                                                                                            Masuda, Esteban
Pardo, Jorge
                                                                                                                                                                      Liao, X.
                                                                                                                                                                                      Li, Congfen
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                                                                                                                                                                        Charlene
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Pred. No. 0.17;
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                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                      SOFTWARE: FO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 98, Application US/10270333
GENERAL INFORMATION:
APPLICANT: CTAVChik, Anibal
TITLE OF INVENTION: ISOLATED G-PROTEIN COUPLED RECEPTORS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING GPCR PROTEINS, AND USES
TITLE OF INVENTION: THEREOF AS INSECTICIDAL TARGETS
FILE REFERENCE: CL000/333CON
                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60/168,677
PRIOR FILING DATE: 1999-12-03
PRIOR APPLICATION NUMBER: 60/175,691
PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR FILING DATE: 2000-03-23
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CURRENT FILING DATE: 2002-10-15
                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -10-270-333-98/c
                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Drosophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/284,763
PRIOR FILING DATE: 2001-04-18
PRIOR APPLICATION NUMBER: US 09/971,228
PRIOR FILING DATE: 2001-10-03
NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Ver. 2.1
        1161
                                                                        1221 CGAATAGCGGCACAGCAGGACCTTCTTGAACGCCTGCCGGAACACCTTGTTGAAGATGGT 1162
                                                                                                                                                                                                                                                                                                LENGTH: 1251
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NAME/KEY: CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: mutant #3, C-terminally truncated variant of OTHER INFORMATION: human endothelial differentiation G-protein OTHER INFORMATION: coupled receptor (GPCR) 1 (EDG1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1333 CGGGCACTTGCAGCAGGACATGATCCGGATGAAGGCCCGACGCATCTCCTTGTTGGTCAG 1274
                                     184 GTACAGGAGCGGGTTGATGACCGAGCTGAGGTAG 217
                                                                                                             124 CGACAGGCGGCAGCACCACCTGCACGAACACCCGGCGAAACTGCTGCGAGGACACCGT 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1273 AGTGTAAATGATGGGGTTGGTGCCGGAGTTGAG 1241
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                                                                                                                                                    Local 59;
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GTAGAAGATGGGGTTCACCATGGAGCTAGCGTAG 1128
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                                                                                                                                                                            62.8%;
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                                                                                                                                                                          Score 38; DB 6; Length 1251; Pred. No. 0.26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 38.6; DB 1;
Pred. No. 0.19;
                                                                                                                                                          Mismatches
                                                                                                                                                      35; Indels
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US-10-270-333-97/c

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SEQ ID NO 115
LENGTH: 5927
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                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/168,677
PRIOR FILING DATE: 1999-12-03
PRIOR APPLICATION NUMBER: 60/175,691
PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR FILING DATE: 2000-03-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Cravchik, Anibal
TITLE OF INVENTION: ISOLATED G-PROTEIN COUPLED RECEPTORS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING GPCR PROTEINS, AND USES
TITLE OF INVENTION: THEREOF AS INSECTICIDAL TARGETS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/270,333
CURRENT FILING DATE: 2002-10-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS:
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NUMBER OF SEQ ID NOS: 198
SOFTWARE: FastSEQ for Windows Version 4.0
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TITLE OF INVENTION: ISOLATED G-PROTEIN COUPLED RECEPTORS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING GPCR PROTEINS, AND USES
TITLE OF INVENTION: THEREOF AS INSECTICIDAL TARGETS
ETLE REFERENCE: CL000733CON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/270,333
CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/168,677
PRIOR FILING DATE: 1999-12-03
PRIOR APPLICATION NUMBER: 60/175,691
PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR FILING DATE: 2000-01-12
PRIOR APPLICATION NUMBER: 60/191,638
                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
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TYPE: DNA
ORGANISM: Drosophila
1431 ACACGAGCTGCTGGAGGCCAGAGTCTGGGCGGTGACCTGGATGACTAGCTTGGTGAGGGT 1490
                                                                                            1371 CACCTTGTAGAAGGCCTTCCGGAAATTCTCGGAGAGGAAGGCGTAGAGCAGCAGCGATTGAT 1430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8178 GTAGAAGATGGGGTTCACCATGGAGCTAGCGTAG 8145
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                                          202 GACCGAGCTGAGGTAGAAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAA 261
                                                                                                                                         142 CACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGAT 201
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                                                                                                                                                                                                                            Local Similarity
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55.9%;
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                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                               Score 37.4; DB 6; Length 5927; Pred. No. 0.79;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 38; DB 6;
Pred. No. 0.67;
0; Mismatches 3
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                                                                                                                                                                                              Indels
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APPLICANT: He, M

APPLICANT: Sheldon, P

TITLE OF INVENTION: Mitomycin biosynthetic gene cluster
FILE REFERENCE: 600.456US1

CURRENT FILING DATE: 2002-10-09
PRIOR APPLICATION NUMBER: US 09/266,965
PRIOR APPLICATION NUMBER: US 09/264,447
PRIOR FILING DATE: 1996-03-12
PRIOR FILING DATE: 1996-08-19
PRIOR APPLICATION NUMBER: US 08/624,447
PRIOR APPLICATION NUMBER: US 08/624,447
PRIOR FILING DATE: 1996-08-19
PRIOR APPLICATION NUMBER: US 08/624,363
PRIOR APPLICATION NUMBER: US 08/133,963
PRIOR APPLICATION NUMBER: US 08/133,963
PRIOR FILING DATE: 1993-10-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: ISOLATED G-PROTEIN COUPLED RECEPTORS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING GPCR PRITITLE OF INVENTION: THEREOF AS INSECTICIDAL TARGETS
FILE REFERENCE: CL000733CON
CURRENT APPLICATION NUMBER: US/10/270,333
CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/168,677
PRIOR FILING DATE: 1999-12-03
PRIOR APPLICATION NUMBER: 60/193,691
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR APPLICATION NUMBER: 60/191,638
PRIOR FILING DATE: 2000-03-23
NUMBER: OF SEQ ID NOS: 198
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US-10-270-333-116
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                        SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 36 LENGTH: 852
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 36, Application US/10267255 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 5.8%;
Best Local Similarity 54.5%;
Matches 73; Conservative
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SEQ ID NO 116
LENGTH: 990
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GENERAL INFORMATION:
APPLICANT: Cravchik, Anibal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -10-267-255-36/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Sherman, D
APPLICANT: Mao, Y
APPLICANT: Varoglu, N
                                                                                                        NUMBER OF SEQ ID NOS:
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    135 AGCACAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCG 194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             868 TGAGGGTGTTCGTC 855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              195 GGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACG 254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              255 CCCGGAAGTAGGAC 268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                928 GATTGATACACGAGCTGCTGTAGGCCAGAGTCTGGGCGGTGACCTGGATGACTAGCTTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     988 AGAAATTCGCCTTGTAGAAGGCCTTCCGGAAATTCTCGGAGAGGAAGGCGTAGAGCAGCG 929
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             262 GTAGGAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Varoglu, M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 36.4; DB 6; Length 990; Pred. No. 0.63;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61; Indels
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APPLICANT: Varoglu, M
APPLICANT: He, M
APPLICANT: Sheldon, P
ITILE OF INVENTION: Mitomycin biosynthetic gene cluster
FILE REFERENCE: 600.456US1
CURRENT APPLICATION NUMBER: US/10/267,255
CURRENT FILING DATE: 2002-10-09
PRIOR APPLICATION NUMBER: US 09/266,965
PRIOR APPLICATION NUMBER: US 09/264,447
PRIOR APPLICATION NUMBER: US 08/624,447
PRIOR APPLICATION NUMBER: US 08/624,447
PRIOR APPLICATION NUMBER: DS 08/624,447
PRIOR APPLICATION NUMBER: PCT/US94/11279
PRIOR APPLICATION NUMBER: PCT/US94/11279
PRIOR APPLICATION NUMBER: PCT/US94/11279
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; ORGANISM: Streptomyces lavendulae
US-10-267-255-76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 76
LENGTH: 53500
                                                                                                                                                                                                                                                                                                                                                                        Query Match 5.8%;
Best Local Similarity 47.9%;
Matches 104; Conservative
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 08/133,963 PRIOR FILING DATE: 1993-10-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Sherman, D
APPLICANT: Mao, Y
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                        7046 CGCTCGAAGAAGTCGGTGAGCACCAGACGGCCCCCGGGCCGAGCACCCGGCCGACCTGG
7226 GCGAGCCCGGCCTCCTCGGCCAGCGCGTTGGCCCGGA 7262
                                                                                                                                                                                7106 GCGAGCACCTGGGCGCGGTCGGGCATGTGGATGATCGATTCGAGGGCGATGACGGCGTCG
                                                                                        7166 AAGCTCTCGTCCTCGAAGGGGAGGTCCATCGCGTCGGCCCGCTGGAAGCGCCCCGGTCG
                                                                                                                   218 AAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAG
                                         278 TCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAA 314
                                                                                                                                                                                                                            158 CGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    359 GCGAGCCCGGCCTCCTCGGCCAGCGCGTTGGCCCCGGA 323
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            419 AAGCTCTCGTCCTCGAAGGGGAGGTCCATCGCGTCGGCCCGCTGGAAGCGCGCCCGGTCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          479 GCGAGCACCTGGGCGCGGCGTCGGCATGTGGATGATCGATTCGAGGGCGATGACGGCGTCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      158 CGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       539 CGCTCGAAGAAGTCGGTGAGCACCAGACGGCCCCCGGGCCGCAGCACCTGG
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                                                                                                                                                                                                                                                                                                                       98 CGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGCGCAGCACCAGCACCTGCACGAACACC 157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98 CGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGCGCGGCAGCACCAGCACCTGCACGAACACC 157
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47.9%;
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RESULT 14 PCT-US02-32094-2/c

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RESULT 15

US-10-265-872-2/c

US-10-265-872-2/c

Sequence 2, Application US/10265872

Sequence 2, Application US/10265872

GENERAL INFORMATION:

APPLICANT: Williamson, Mark

TITLE OF INVENTION: COUPLED RECEPTOR

TITLE OF INVENTION: COUPLED RECEPTOR

FILE REFERENCE: MPIO1-233P1RM

CURRENT APPLICATION NUMBER: US/10/265,872

CURRENT FILING DATE: 2002-10-07

PRIOR APPLICATION NUMBER: 66/328,280

PRIOR FILING DATE: 2001-10-09

NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                      ; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 2
; LENGTH: 1285
; TYPE: DNA
; ORGANISM: Homo sapien
US-10-265-872-2
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; TYPE: DNA
; ORGANISM: Homo sapien
PCT-US02-32094-2
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Best Local S
Matches 76
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GENERAL INFORMATION:

APPLICANT: Williamson, Mark

TITLE OF INVENTION: METHODS OF USING 241, A HUMAN G-PROTEIN

TITLE OF INVENTION: COUPLED RECEPTOR

FILE REFERENCE: MPI01-233

CURRENT APPLICATION NUMBER: PCT/US02/32094

CURRENT APPLICATION NUMBER: 9002-10-08

PRIOR APPLICATION NUMBER: 60/328,280

PRIOR APPLICATION NUMBER: 60/328,280

PRIOR FILING DATE: 2001-10-09

NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 2
                                                                                                                                                                                                                                  Query Match
Best Local :
                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1035 GACGGGGTTGGCACAGCTGTTGGCGTAGGAGAGGATGACCACGAAGAAGTAGAGGCCGGC 976
                                                                                                                                       1035 GACGGGGTTGGCACAGCTGTTGGCGTAGGAGAGGATGACCACGAAGAAGTAGAGGCCCGC 976
975
                                                                                                         190
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                                 250
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Local Similarity 53.1%;
nes 76; Conservative (
                                                                                                                                                                                                                  Local Similarity
les 76; Conserv
GGAGGCGGCTCCTGGGGCAGCG 953
                                GTACGCCCGGAAGTAGGACCTCG 272
                                                                                                       GAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCAT 249
                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                  5.7%; Score 35.8; 53.1%; Pred. No. 1;
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; Pred. No. 1;
0; Mismatches
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Search completed: November 7, 2002, 17:57:05 Job time :  $38.4612 \ \text{secs}$ 

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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Perfect score:
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Gapop 10.0 , Gapext 1.0
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em_gss_fun:*
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                AI936826
AI990500
                                        BQ006545
BM669397
BM547680
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BQ006545 UI-H-EI1-
BM669397 UI-E-DW1-
BM547680 AGENCOURT
AI936826 Wp69h10.x
AI990500 ws40d07.x
AW590950 hg51e12.x
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## ALIGNMENTS

JOURNAL COMMENT REFERENCE AUTHORS TITLE VERSION KEYWORDS SOURCE ORGANISM RESULT 1 BQ006545/c FEATURES ACCESSION DEFINITION source Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Catarrhini; Hominidae; Homo.

1 (bases 1 to 702)

1 (bases 1 to 702)

2 1 (bases 1 to 702)

2 NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

2 NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

2 National Cancer Institute, Cancer Genome Anatomy Project (CGAP),

2 Tumor Gene Index

2 Unpublished (1997)

3 Contact: Robert Strausberg, Ph.D.

5 Email: cgapbs-remail.nih.gov

7 Iissue Procurement: Dr. Jose Mercuende

5 CDNA Library preparation: Dr. M. Bento Soares, University of Iowa

6 CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa

6 CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa

6 Clone Distribution: Clone distribution information can be found

7 through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov

8 POLYA-Yes. BQ006545 702 bp mRNA linear EST 26-MAR-UI-H-EI1-aza-n-21-0-UI.sl NCI\_CGAP\_EII Homo sapiens cDNA clone IMAGE:5846228 3', mRNA sequence.
BQ006545 EST BQ006545.1 GI:19731445 Location/Qualifiers EST 26-MAR-2002

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ACAGTGTAAAAAAAAAAAAAAAAAA

EST 27-FEB-2002

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                                                                      TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT
                                                                                                                                                                                                                                                                                                                                                                       /lab_host="DHIOB (Life Technologies)"
//lab_host="DHIOB (Life Technologies)"
//note="organ: Left Pelvis; Vector: pT7T3-Pac (Pharmacia)
with a modified polylinker; Site_1: EcoR I; Site_2: Not I;
with a modified polylinker; Site_1: EcoR I; Site_2: Not I;
with a modified polylinker; Site_1: Scor I; Site_2: Not I;
close="organized cond. Ibrary containing the following tissue(s): Chondrosarcoma. The library was constructed according to Bonaldo, Lennon and Soares, Genome Research, 6:791-806, 1996. First strand cDNA synthesis was primed with an oligo-dT primer containing a Not I site. Double stranded cDNA was ligated to an EcoR I
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/db_xref="taxon:9606"
/clone="IMAGE:5846228"
/clone_lib="NCI_CGAP_EII"
/tissue_type="chondrosarcoma"
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TAG_SEQ-ACACTTGCAC"
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                                                                                                                                           Query Match
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  625
                                                                                                                       Local Similarity
                                             1 AGTTCTCCTTGCAGAGGACTGGCGCGGGGCGGACGCGAAGAGCAAACGGGCGCTGCACAAAGCG 60
UI-E-DW1-ahd-c-11-0-UI.S1 UI-E-DW1 Homo sapiens cDNA clone UI-E-DW1-ahd-c-11-0-UI 3', mRNA sequence.
BM669397
BM669397.1 GI:18979294
EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: msoares@blue.weeg.uiowa.edu
Tissue Procurement: Dr. Gregg Hageman
cDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa
DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
Clone Distribution: Researchers may obtain clones from Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         University of Iowa
451 Eckstein Medical Research Building Iowa City,
Tel: 319 335 8250
Fax: 319 335 9565
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Bonaldo, M.F., Lennon, G. and Soares, M.B.
Normalization and subtraction: two appr
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                Conservative
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//lab_bost="mbl08" (Life Technologies) (Ti phage resistant)"
//note="Organ: eye; Vector: pT7T3-Pac (pharmacia) with a
modified polylinker; Site_1: Ecot I; Site_2: Not I;
UI-E-NMI is a normalized cDNA library containing the
following tissue(s): lens. The library was constructed
according to Bonaldo, Lennon and Soares, Genome Research,
6:791-806, 1996. First strand cDNA synthesis was primed
with an oligo-dT primer containing a Not I site. Double
stranded cDNA was ligated to an EcoR I adaptor, digested
with Not I, and cloned directionally into pT7T3-Pac
vector. The oligonucleotide used to prime the synthesis of
first-strand cDNA contains a library tag sequence that is
located between the Not I site and the (dT)18 tail. The
sequence tag for this library is CGATTACCGA. This library
was created for the program, Gene Discovery in the Visual
System, supported by National Eye Institute (NEI).
TAG_LIB-UI-E-DWI
TAG_SEQ-CGATTAGCGA"
183 t 1 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="UI E-DW1 and -c-11-0-UI"
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/db_xref="taxon:9606"
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facilitate gene

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                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Invitrogen
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Bioscience Corporation
Clone distribution: MGC Clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM12722 row: h column: 07
                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 696)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                      AGENCOURT_6507108 NIH_MGC_124
5', mRNA sequence.
                                                                                                                                                                      Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
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                                 quality sequence stop: 641.
Location/Qualifiers
/organism="Homo sapiens"
/db_xref="taxon:9606"
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4 Homo
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RESULT 4 \AI936826/c

AI936826

ACCESSION VERSION

GPR39. ; AI936826

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AI936826.1 EST.

AI336826 678 bp mRNA linear EST 08-MAR-2000 wp69hl0.xl NCI\_CGAP\_Brn25 Homo sapiens cDNA clone IMAGE:2467075 3' similar to SW:GP39\_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR

KEYWORDS

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ACAGTGTAAAAAANNNNNAA 685
                                               TAAATGAATAAAACCATAAAATATTTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="IMAGE:5727798"
/clone_lib="NIH_MGC_124"
/tissue_type="hippocampus"
/lab_host="DH10B"
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98.4%;
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Pred. No. 1.6e-100;
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National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index Unpublished (1998)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /tissue_type="anaplastic oligodendroglioma"
/lab_host="DH10B"
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/clone_lib="NCI_CGAP_Brn25"
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/db_xref="taxon:9606"
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99.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      found through the I.M.A.G.E. Consortium/LLNL www-bio.llnl.gov/bbrp/image/image.html Insert Length: 1212 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert Strausberg, Ph.D.
Email: Cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D.,
R. Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AI990500 616 bp mRNA linear EST 08-MAR-2000 ws40d07.x1 NCI_CGAP_GC6 Homo sapiens cDNA clone IMAGE:2499661 3' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
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National Cancer Institute, Cancer Genome Ana
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DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can b
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/note="Vector: pf7773D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Plasmid DNA from the normalized library NCI_GCAP_GC4 was prepared, and scircles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1257096-1258631, 1459064-1470983, and 1475592-1476743). Subtraction by Bento Soares and M. Fatima Bonaldo. " a 182 c 147 g 167 t
                                                                                                                                                                                                                            /clone_lib="NCI_CGAP_GC6"
/tissue_type="pooled germ cell tumors"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                /clone="IMAGE:2499661"
                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
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Score 601.2;

DB 9;

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                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                      AW590950
649 bp mRNA linear EST 22-hg51e12.x1 NCI_CGAP_GC6 Homo sapiens cDNA clone IMAGE:29491 similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED
                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher A.
R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: M. Bento
                                                                                                     Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 649)
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Catarrhini; Hominidae
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found through the I.M.A.G.E. Consortium/LLNL at:
image.llnl.gov/image/html/iresources.shtml
                                                                  DNA Sequencing by: Washington University Genome Clone distribution: NCI-CGAP clone distribution
                                                                     information
                                                                                               Sequencing Center
                                                                     can be
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Possible reversed clone: similarity on wrong strand primer: -40UP from quality sequence stop: Gibco 457

132 ø /note="vector: pT7T3D-Pac (Pharmacia) with a modified polylinker: Site\_1: Not I; Site\_2: Eco RI; Plasmid DNA from the normalized library NCI\_CGAP\_GC4 was prepared, and scircles were made in vitro. Following HAP purification this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library (cloneIDs 1257096-1258631, 1469064-1470983, and 1475592-1476743). Subtraction by Bento Soares and M. Fatima Bonaldo. " 189 c 155 g 173 t /tissue\_type="pooled germ
/lab\_host="DH10B" /clone\_lib="NCI\_CGAP\_GC6" /db\_xref="taxon:9606" /clone="IMAGE:2949166" /organism="Homo sapiens" location/Qualifiers cell tumors"

Similarity Conservative 96.2%; 99.5%; 0; Score 601.2; Pred. No. 7.3e 0; Mismatches .3e-99 DB 10; ω --Indels Length 0 Gaps 0

547 GGCGCTGTCGGTGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG 1 AGTTCTCCTTGCAGAGGACTGGCGCCGGGACGCGAAGAGCAACGGGCGCTGCACAAAGCG 60 CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG CGTGTACAGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAG CAGCGACAGGCGCAGCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACAC CAGCGACAGGCGGCACACAGCACCTGCACGAACACCCGGCCGAAACTGCTGCGAGGACAC GGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTG AGTTCTCCTTGCAGAGGACTGGCGCCCGGTACGCGAAGATCAACGGGCGCTGCACAAAGCG 180 240 428 488 120 548 368

GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC GAGGATCATGTACGCCCGGAAGTAGGACCTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGC 308 300

CATGATCCTCCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC CATGATCCTCCGAATCTGGGTTGGGCATCCAGCATACGGCCAATGTCACAACAATCAGCCC

420

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TGGGCAGACACGAGCAGGAGGAGAGACAGAAAAAGAAAAACACAGCATGAGAACACAG

TAAATGAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 480 TAAATAAATAAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAAT 128

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KEYWORDS
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                                                                                        AGGCGGCAGCACCACCTGCACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTAC
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      AGGAGCGGGTTGATGACCGAGCTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Seq primer: -40UP from Gibco
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Location/Qualifiers
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202 c 1
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/clone_lib="Soares_NSF_F8_9W_OT_PA_P_S1"
/lab_host="DH10B"
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/db_xref="taxon:9606"
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Pred. No. 8.3e-99;
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                                                                                                                                                                                                                                                                                                                                          Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Burt Feuerstein, M.D., Mark Israel, M.D.
cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute. Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo (Dases 1 to 690)
NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute / National Institute of Neurola Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1998)
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                                                                                                                                                                                                                                                                                                                          www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                        quality sequence stop: 402
/clone="IMAGE:2620569"
/clone_lib="NCI_CGAP_Brn50"
/clone_lib="NCI_CGAP_Brn50"
/liss_ne_type="medulloblastoma"
/lab_host="DH10B (phage resistant)"
/note="Organ: brain; Vector: pT'773D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from medulloblastoma tumor tissue, and was then primed with a
                                                                                                                                                    /db_xref="taxon:9606"
                                                                                                                                                                            /organism="Homo sapiens"
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on wrong

Neurological

Euteleostomi; Homo.

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RESULT 9
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eukaryota; Metazoa; Chordata; Catarrhini; Hominidae 1 (bases 1 to 645)
NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.National Cancer Institute / National Institute of National Cancer Institute / National Institute of National Cancer Institute / National Institute of National Cancer Institute
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nac53g11 x1 NCI_CGAP_Brn23 Homo similar to SW:GP39_HUMAN 043194 GPR39. , mRNA sequence.

BF432379
                                                                               Homo sapiens
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Bonaldo, Ph.D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N. Louis, M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1998)
Contact: Robert Strausberg,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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DNA Sequencing by: Washington University Senome Sequencing C

Clone distribution: NCI-CGAP clone distribution information

Cound through the I.M.A.G.E. Consortium/LLNL, send email to:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer: -40UP from Gibco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="NCI_CGAP_Brn23"
/tissue_type="glioblastoma
/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
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98.0%;
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Tumor Gene Index
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Pred. No. 9.9e-96;
0; Mismatches 12
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Best Local Similarity
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AGCGGGCGCTGTCGGTGGAGTGCGCATGTACGCGCAGGCGCTTCTCGTGGTTGGCGT 492
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National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index Unpublished (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BE858216 551 bp mRNA linear EST 29-SEP-7919907.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:3306972 similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BE858216
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             High quality sequence stop: 490.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 found through the I.M.A.G.E. Consortium/LLNL, send email to:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cDNA Library Preparation: M. Bento Soares, Ph.D.,
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                                                                                                        Conservative
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                                                                                                                                                                                            /db_xref="taxon:9606"
/clone="IMAGE:3306972"
/clone_lib="NCI_CGAP_Brn23"
/tissue_type="glioblastoma (pooled)"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                            'organism="Homo sapiens"
                                                                                                                        87.7%;
99.8%;
                                                                                                  Score 548.4; DB 12,
Pred. No. 2.5e-89;
                                                                                                                                       DB 12;
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nac80b12.x1 NCI_CGAP_Brn23 Homo
similar to SW:GP39_HUMAN 043194
                                                                                                                                                                                                                                                                            (CGAP/BTGAP), Tumor Gene Index
Unpublished (1998)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
                                                                                                                                     CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL, send email to:
                                                                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 540)
NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
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                                                                                                                                                                                                              Bonaldo,
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Disorders and Stroke, Brain Tumor Genome Anatomy Project
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Mammalia; Eutheria; Primațes; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                Tissue Procurement: David N. Louis, M.D.,
                                                                                                                     info@image.llnl.gov
                                                                                                                                                                                                                           cDNA Library Preparation: M. Bento Soares, Ph.D.,
                                                                                                      primer: -40UP from Gibco
                                                                                    quality sequence stop:
            /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                 Location/Qualifiers
                                                                                     467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mRNA linear EST 22-JAN-2001 sapiens cDNA clone IMAGE:3440591 3' PUTATIVE G PROTEIN-COUPLED RECEPTOR
                                                                                                                                                                                                                                                                 Myrna R.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 537)
NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.National Cancer Institute / National Institute of Ne Disorders and Stroke, Brain Tumor Genome Anatomy Pro
                                                                                                                                          GPR39. ;, mRNA sequence.
BF594242
BF594242.1 GI:11686566
                                                                                                                                                                                   BF594242 537 bp mRNA linear EST 12-DEC-2000 7n10d03.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:3564316 3' similar to SW:GP39_HUMAN 043194 PUTATIVE G PROTEIN-COUPLED RECEPTOR
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                                                                                            Homo sapiens
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Pred. No. 1.6e
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No. 1.6e-87;
Genome Anatomy Pro-
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                                                            TTTTCAGTGTTGGACTGACAGCTTCTTTTGCCACAGCAAGAAGAAGAATTTAACACTGTT 549
                                                                                                                                                                                                                                                   CCGAATCTGGTTGGGCATCCAGCATACGGCCAATGTCACCAACAACCAGCCCTGGGCAGAC
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             TCAAACCCGGGGGAGTTGGCTGTGTTAAAGAAAGACCATTAAATGCTTTAGACAGTG
                                            AAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAAT 118
                                                                                                                         AAAACCATAAAATATTTAGCCCCTCTGTTCTGTGCTTACTGGCCAGGAAATGGTACCAAT 489
                                                                                                                                                                       ACGAGCAGGAGGAGAGACAGAGAAAAAGAAAAACACAGCATGAGAACACAGTAAATGAAT 178
                                                                                                                                                                                                     ACGAGCAGGAGGAGAGAGAAAAGAAAACACAGCATGAGAACACAGTAAATGAAT 429
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Contact: Robert Strausberg, |
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seq primer: -40UP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can by
found through the I.M.A.G.E. Consortium/LLNL, send email to:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bonaldo,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CGAP/BTGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cDNA Library Preparation: M. Bento
sonaldo, Ph.D.
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Location/Qualifiers
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/clone_lib="NCI_CGAP_Brn23"
/tissue_type="glioblastoma (pooled)"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
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Pred. No. 2.8e-87;
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  538
                    88 TACGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGCAGCAGCACCAGCACCTG 147
  Email: msoares@blue.weeg.ulowa.edu
Tissue Procurement: Dr. Gregg Hageman
cDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iowa
cDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa
cDNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
Clone Distribution: Researchers may obtain clones from Research
Genetics (www.resgen.com).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BM667957 538 bp mRNA linear EST 27-FEB-2002 UI-E-DW0-agm-j-01-0-UI.S1 UI-E-DW0 Homo sapiens cDNA clone UI-E-DW0-agm-j-01-0-UI 3', mRNA sequence.

BM667957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Program for Rat Gene Discovery and Mapping
University of Iowa
451 Eckstein Medical Research Building Iow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            451 Eckstein Medical Research Building Iowa
Tel: 319 335 8250
Fax: 319 335 9565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 538)
Bonaldo, M.F., Lenno
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                                                                                                    Similarity
                                                                                                                                                                                   104
                                                                                 Conservative
                                                                                                                                                                                                                                     tissue(s): lens. The library was constructed according to Bonaldo, Lennon and Soares, Genome Research, 6:791-806, 1996. First strand cDNA synthesis was primed with an oligo-dT primer containing a Not I site. Double stranded cDNA was ligated to an EcoR I adaptor, digested with Not I, and cloned directionally into pT773-Pac vector. The oligonucleotide used to prime the synthesis of first-strand cDNA contains a library tag sequence that is located between the Not I site and the (dT)18 tail. The sequence tag for the program, Gene Discovery in the Visual System, supported by National Eye Institute (NEI).
                                                                                                                                                                               TAG_TISSUE-human lens
TAG_SEQ-CGATTAGCGA"
144 c 122 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /lab_host="DH10B (Life Technologies) (Tl phage resistant)" /note="Organ: eye; Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site_1: EcoR I; Site_2: Not I; UI-E-DW0 is a cDNA library containing the following
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /db_xref="taxon:9606"
/clone="UI-E-DWO-agm-j-01-0-UI"
/clone_lib="UI-E-DWO"
/tissue_type="lens"
/dev_stage="adult"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
                                                                                               85.2%;
99.3%;
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Pred.
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No. 1.6
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AI018769
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ov32e04.x1 Soares_testis_NHT Homo
3' similar to SW:NTR1_RAT P20789
                                                                                                                                                                                                                DNA Sequencing by: Washington University Genome Sequencing C Clone distribution: NCI-CGAR clone distribution information found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 1394 Std Error: 0.00
                                                                                                                                                                                                                                                                                                                                                            Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: ggapbs-r@mail.nih.gov
CDNA Library Preparation: M. Bento Soares, Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhin1; Hominidae; Homo. 1 (bases 1 to 537)
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National Cancer Institute, Cancer Genome Anat
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                                                                                                                                                            primer: -40m13 fwd. ET from Amersham
h quality sequence stop: 452.
Location/Qualifiers
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                                                        /clone_lib="Soares_testis_NHT"
/note="Vector: pT7T3D-Pac (Pharmacia) with
                      /lab_host="DH10B"
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bp mrNA linear EST 27-AUG-1998
Homo sapiens cDNA clone IMAGE:1639038
O789 NEUROTENSIN RECEPTOR TYPE 1;

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Anatomy Project (CGAP),

M. Fatima Bonaldo

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                                                                                                                                                                         x939C02.xl NCI_CGAP_Utl Homo sapiens cDNA clone IMAGE:2629922 3/ similar to SW:GP39_HUMAN O43194 PUTATIVE G PROTEIN-COUPLED RECEPTOR GPR39. ; mRNA sequence.
           Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg,
                                                                              Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 561)
                                                                                                                                     human.
                                                                                                                                                  EST
Email: cgapbs-r@mail.nih.gov
                                                                                                                        Homo
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                                                 National Cancer Institute, Cancer
                                                                 NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
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Emmert-Buck, M.D., Ph.D.
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Site_2: NotI; Cloned unidirectionally. Primer: Oligo
Average insert size 1.75 kb. Life Technologies catalog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="IMAGE:2629922"
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/tissue_type="well-differentiated endometrial
adenocarcinoma, 7 pooled tumors"
/lab_host="PH10B"
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/db_xref="taxon:9606"
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GenCore version 5.1.3
Copyright (c) 1993 - 2002 Compus
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Database : OM nucleic -Minimum DB seq length: 0
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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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| ull inser<br>or cortex or<br>thordata; or                      | 1832 b<br>37182 fis                      | ALIGNMENTS | AF514845<br>BC034500<br>AF044209<br>AC005250 | AF176571<br>AX056994 | AC069297<br>AL133395 | AC110232 | AC074245<br>AC074389 | AC126321<br>AC027118 | AL645938 | AC112072 | AY094780<br>AL645986 | AKU26521<br>BC028978 | AY118558<br>AX285020 | AF082874 | AC123338 | AB041649         | AC010974 | AX136556   | AX093381      | AX136698     | AF034633 | BC011449 | AX136281  | AX319944 | AX358762 | AK094501<br>AC079773 | ID                         | SUMMARIES |
| t sequence).<br>cDNA to mRNA<br>Craniata; Ve;<br>Catarrhini; I | bp mRNA<br>is, clone BR<br>GTPase-acti   | ENTS       |                                              |                      |                      |          |                      |                      |          |          |                      |                      |                      |          |          |                  |          |            |               |              |          |          |           |          |          |                      |                            | IES       |
| NA, clo<br>Vertebr                                             | nRNA line<br>ne BRALZ2001<br>-activating |            |                                              |                      |                      |          |                      |                      |          |          |                      |                      |                      |          |          |                  |          |            |               |              |          |          |           |          |          |                      |                            |           |
| ne_li<br>ata;<br>idae;                                         | prot                                     |            | AF514845<br>BC034500<br>AF044209<br>AC005250 | AF1765               | AC0692               | AC1102   | AC0742               | AC1263<br>AC0271     | AL60/8   | AC1120   | AY0947               | BC0289               | AY1185<br>AX2850     | AF0828   | AC1233   | AC1 244<br>AB041 | AC0109   | AX0931     | AX1301        | AX1366       | AF0346   | BC0114   | AX1362    | AX3199   | AX3587   | AK0945<br>AC0797     | Descri                     |           |
| ):BRALZ<br>Juteleo                                             | PRI 15-J<br>weakly s<br>ein (RGS2        |            |                                              | 44 Zed<br>71 Alt     |                      |          |                      | ~ ~                  | ω,       |          |                      | 78 Homo              | 58 Dr                | 74 Medi  |          |                  |          |            |               |              |          | 49 Homo  |           | S O      |          | <b>#</b> #           | ption                      |           |
| b:BRALZ2<br>Euteleostom1;<br>Homo.                             | UL-200<br>imilar                         |            | na cate<br>no sapi<br>no sapi<br>no sapi     | ternari              |                      | = =      | no sapi<br>no sapi   | ٥×                   |          | 00 (     | Drosophil            | no sapi              |                      | dicago   | on st    | us muscu         | no sapi  | Sequence   | quence        | 2            | mo sapi  | 90       | enc       | equence  | ence     | Sa                   | 1<br>1<br>4<br>2<br>5<br>6 |           |
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| 181 C<br>415 C<br>241 C<br>475 C<br>301 C<br>361 C<br>361 C            | ery Match st Local S1 tches 1597; 1 GGCAA               235 GGCAA 61 GTGTG             295 GTGTGA 121 GGTGA                                               | CDS                                                                                                                                                                                                                                                                                                                                                                                                                                                       | ENT URES Source                                                                                                                                                                                                                                                                                                                                                                                    | THORS TLE URNAL RENCE THORS TLE URNAL                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |
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| CCGCAAGTCCTGTGCATCATCAGCGGCCTGTCTCATCGCCTCTGCCGGGTACCAGTCCTT 240       | ery Match 88.1%; Score 1427; DB 9; Length 1832; st Local Similarity 99.9%; Pred. No. 0; 1 GGCAACTTTTGCCGGATTGTTCTTGCTTCCAGGCTTTGCGCTGCAAATCCAGTGCTACCA 60 | /organism="Homo sapiens" /db_xref="taxon:9606" /clone="BRALIZ001550" /tissue_type="alzheimer cortex" /clone_lib="BRALIZ2" /note="cloning vector: pME18SFL3" /15640 /note="unnamed protein product" /codon_Start=1 /protein_id="BAC04368.1" /db_xref="GI:2175576" /translation="MWVLGIANATECGLFLLPGFALQIQCYQCEEFQLNNDCSSPEFIV NCTVNVQDMCQKEVMEDSAGINYRKSCASSAACILIASAGYQSFCSPGKLNSVCISCCN TPLCNGPRPKKRGSSASALRPGLRTTILFLKLALFSAHC" 411 a 501 c 513 g 407 t | (E-mail:genomicsehri.co.jp, Tel:<br>(E-mail:genomicsehri.co.jp, Tel:<br>NEDO human cDNA sequencing proje<br>Economy, Trade and Industry of J<br>Research Association for Biotech<br>construction: Helix Research Ins<br>Key Technology Center etc.); 5'-<br>HRI, and Biotechnology Center, N<br>Evaluation; clone selection for<br>RAB; annotation: HRI and RAB.<br>Location/Qualifiers<br>1. 1832 | Kawakami, B., Sugiyama, A., Takemoto, M., Sugiyama, T., Irie, R., Otsuki, T., Sato, H., Wakamatsu, A., Ishii, S., Yamamoto, J., Isono, Y., Kawai-Hio, Y., Saito, K., Nishikawa, T., Kimura, K., Yamashita, H., Matsuo, K., Nakamura, Y., Sekine, M., Kikuchi, H., Kanda, K., Wagatsuma, M., Murakawa, K., Kanehori, K., Takahashi, Fujii, A., Oshima, A., Suzuki, Y., Sugano, S., Nagahari, K., Masuho, Y., Nagai, K. and Isogai, T. NEDO human cDNA sequencing project Unpublished ( hases I to 1832) Isogai, T. and Yamamoto, J. Direct Submission Submitted (04-JUL-2002) Takao Isogai, FLJ Project(HRI Team); 2-6-7 Kasusa-Kanatari, Kiaararu, Chiha 292-0812, Japan |
| 04<br>04<br>05<br>05<br>05                                             | 0                                                                                                                                                         | OY OY                                                                                                                                                                                                                                                                                                                                                                                                                                                     | 0 00 00 00 00 00 00 00 00 00 00 00 00 0                                                                                                                                                                                                                                                                                                                                                            | 0                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        |
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Submitted (09-TAN-2002) Department of Genetics, Washington Submitted (09-TAN-2002) Department of Genetics, Washington University, 4444 Forest Park Avenue, St. Louis, Missouri 63108, University, 4444 Forest Park Avenue, St. Louis, Missouri 63108, On Aug 9, 2001 this sequence version replaced gi:14488388.

On Aug 9, 2001 this sequence Tananaity Genome Sequencing Center
                                                                                                                    This sequence was finished as follows unless otherwise noted: all regions were double stranded, sequenced with an alternate chemistry, or covered by high quality data (i.e., phred quality >= 30); an attempt was made to resolve all sequencing problems, such as compressions and repeats; all regions were covered by sequence from more than one subclone; and the assembly was confirmed by
                                                                                                                                                                                                                                                                                                                      NOTICE: This sequence may not represent the entire insert of this clone. It may be shorter because we only sequence overlapping clone sections once, or longer because we provide a small overlap
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Waterston, R.
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Homo sapiens BAC clone RP11-258B17
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Mapping information for this clone was provided by Dr. John D McPherson, Department of Genetics, Washington University, St.
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Contact: sapiens@watson.wustl.edu
------ Summary Statistics
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VECTOR: pBACe3.6
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| AAGTO                                                        | 80.9<br>99.9<br>tive                         | 13<br>_fami                     | 63            | 93    |         | _ ramı              | 73      | 3. ami    | _ramı   | 4 · . 3       | 83            | 3.3          | 7 3          | 41            | 4 · · · · 2 | 2          | 6 2<br>fami      | 0        | 8 2<br>fami  | fami           | 9 2     | fami         | 6              | 62                                     | 62      |           | 52      | 8 2<br>6 2 |              | 4        | _fam1<br>82  | _fam1     | fami         |
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.C., Gurney,A.L., Smith,V., 335 DB đ 6 0; and Length 1524; Indels linear nucleic acids Euteleostomi; PAT Stephan, J.P., Ното 15-FEB-2002 620 480 600 660 680 420 360 440 740 540 560 500 240 320 180 ۲,

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ORGANISM RESULT 5 AX319944/c LOCUS ACCESSION 밁 20 В 80 밁 ργ 멍 Qy 밁 δÃ 밁 Qy В δÃ В οy DЬ Ωy 망 δõ δÃ Qy Qy REFERENCE 망 В 밁 밁 AUTHORS TITLE JOURNAL 1520 1381 1340 1221 1021 1101 1041 901 981 921 861 801 781 AGAGA AGAGA 1385 CTCGTCCAGTCGTGCTTGGGTTTGGCCGCAGCCATGATCCTCCGAATCTGGGTTGGGCATC CTGAGGTAGAAAAACGTCTCCGAGAAGGGGAGGAGGATCATGTACGCCCGGAAGTAGGAC ACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAG CGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGGCAGCACACGCACCTTGC ACGCGAAGAGCAACGGCGCTGCACAAAGCGGGCGCTGTCGGTGGTGGAGTGCGCATGTA CGCTCTGAAAAGTGCTTAAGAAAATCTTCTCAGTTCTCCTTGCAGAGGACTGGCGCCGGG GTGAGAGTGACGTTTTCTTAGGGCTGGAGGGCCAGTTCCCACTCAAGGCTCCCTCGCTTG TTTTGTAAAGCCTCCAGTGTCCATTCCCTGATGGGGGCATAGTTTGAGACTGCAGA CGGAGTCAAGGCCAGGCTGGCAGAGTCAGTCCTTAGAAGTCACTGAGGTGGGCATCTGCC CGCGCAGGCGCTTCTCGTGGTTGGCGTGCTGCAGCGACAGGCGGCAGCACCACC-TGC ACATTCAAACTTCATGCTCCTGAAAACCATTCTCTGCAGCAGAATTGGCTGGTTTCGCGC ACATTCAAACTTCATGCTCCTGAAAACCATTCTCTGCAGCAGAATTGGCTGGTTTCGCGC TTTTGTAAAGCCTCCAGTGTCCATTCCATCCCTGATGGGGGCATAGTTTGAGACTGCAGA ACGAACACCCGCCGAAACTGCTGCGAGGACACCGTGTACAGGAGCGGGTTGATGACCGAG GTGAGAGTGACGTTTTCTTAGGGCTGGAGGGCCAGTTCCCACTCAAGGCTCCCTCGCTTG AX319944 Sequence AX319944 Galvin,K.A. and Rudolph-Owen,L.A. Methods and compositions for the diagnosis cardiovascular and tumorigenic disease usin Patent: WO 0181634-A 3 01-NOV-2001; Millennium Pharmaceuticals, Inc. (US) Homo Eukaryota; Metazoa; Mammalia; Eutheria; AX319944.1 1524 sapiens ω GI:17901491 2528 bp Patent WO0181634. Chordata;
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